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 Glu Asp Ala Ala Gly Leu Gly Gly Pro Asp Gly Lys Ser Gly Arg Trp  
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 Gln Tyr Glu Ile Glu Thr Asp Pro Ala Leu Thr Tyr Val Glu Gly Val  
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 Cys Val Val Trp Phe Thr Phe Glu Phe Leu Val Arg Ile Val Phe Ser  
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Val Ala Ile Leu Pro Phe Tyr Leu Glu Val Gly Leu Ser Gly Leu Ser  
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370 375 380

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Tyr Tyr Ala Glu Arg Val Gly Ala Gln Pro Asn Asp Pro Ser Ala Ser  
405 410 415

Glu His Thr Gln Phe Lys Asn Ile Pro Ile Gly Phe Trp Trp Ala Val  
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Val Thr Met Thr Thr Leu Gly Tyr Gly Asp Met Tyr Pro Gln Thr Trp  
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Ser Gly Met Leu Val Gly Ala Leu Cys Ala Leu Ala Gly Val Leu Thr  
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Ile Ala Met Pro Val Pro Val Ile Val Asn Asn Phe Gly Met Tyr Tyr  
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Ser Leu Ala Met Ala Lys Gln Lys Leu Pro Arg Lys Arg Lys Lys His  
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Gly Asn Ala Leu Arg Leu Ser Pro Val Thr Ser Pro Tyr Asn Ser Pro  
610 615 620





Val Cys Lys Ser Ser Asp Trp Phe Ile His Thr Cys Met Ala Ala Lys  
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Asp Val Pro Ala Val Ala Glu Glu Phe Met Ser Met Phe Gly Lys Leu  
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Tyr Pro Leu Leu Ala Phe Gly Leu Pro Leu Phe Phe Ala Ser Phe Tyr  
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Phe Trp Arg Ala Tyr Asp Gln Cys Lys Lys Arg Gly Thr Lys Thr Gln  
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Lys Gly Val Trp Lys Trp Met Ile Thr Lys Lys Pro Pro Thr Val Ser  
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Glu Ser Gln Glu Thr Pro Ala Gly Asn Ser Glu Gly Leu Pro Asp Lys  
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Val Pro Ser Pro Glu Ser Pro Ala Ser Ile Pro Glu Lys Glu Lys Pro  
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Ser Ser Pro Ser Ser Gly Lys Gly Lys Thr Glu Lys Ala Glu Ile Pro  
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Ile Leu Pro Asp Val Glu Gln Phe Trp His Glu Arg Asp Thr Val Pro  
370 375 380

Ser Val Gln Asp Asn Asp Pro Ile Pro Leu Gly Thr  
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 <213> Homo sapiens

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 Ser Gln Thr Arg Gly Ile Ser Ser Leu Pro Arg Ser Tyr Thr Met Asp  
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 Asp Ala Trp Lys Tyr Asn Gly Asp Ile Glu Asp Ile Lys Arg Thr Pro  
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 Val Thr Arg Leu Pro Ser Pro Thr Ser Pro Phe Ser Ser Leu Ser Gln  
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 Asp Gln Ala Ala Thr Ser Lys Ala Thr Leu Ser Ser Thr Ser Gly Leu  
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Glu Glu Lys Gly Ala Thr Tyr Pro Ser Glu Ile Pro Lys Glu Asp Ser  
 50 55 60

Thr Thr Phe Ala Lys Arg Glu Asp Arg Val Thr Thr Glu Ile Gln Leu  
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Pro Ser Gln Ser Pro Val Glu Glu Gln Ser Pro Ala Ser Leu Ser Ser  
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Leu Arg Ser Arg Ser Thr Gln Met Glu Ser Thr Arg Val Ser Ala Ser  
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Leu Pro Arg Ser Tyr Arg Lys Thr Asp Thr Val Arg Leu Thr Ser Val  
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Val Thr Pro Arg Pro Phe Gly Ser Gln Thr Arg Gly Ile Ser Ser Leu  
 130 135 140

Pro Arg Ser Tyr Thr Met Asp Asp Ala Trp Lys Tyr Asn Gly Asp Val  
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Glu Asp Ile Lys Arg Thr Pro Asn Asn Val Val Ser Thr Pro Ala Pro  
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Ser Pro Asp Ala Ser Gln Leu Ala Ser Ser Leu Ser Ser Gln Lys Glu  
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 Val Ala Ala Thr Glu Glu Asp Val Thr Arg Leu Pro Ser Pro Thr Ser  
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 Pro Phe Ser Ser Leu Ser Gln Asp Gln Ala Ala Thr Ser Lys Ala Thr  
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 Leu Ser Ser Thr Ser Gly Leu Asp Leu Met Ser Glu Ser Gly Glu Gly  
 225 230 235 240  
 Glu Ile Ser Pro Gln Arg Glu Val Ser Arg Ser Gln Asp Gln Phe Ser  
 245 250 255  
 Asp Met Arg Ile Ser Ile Asn Gln Thr Pro Gly Lys Ser Leu Asp Phe  
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 Gly Phe Thr Ile Lys Trp Asp Ile Pro Gly Ile Phe Val Ala Ser Val  
 275 280 285  
 Glu Ala Gly Ser Pro Ala Glu Phe Ser Gln Leu Gln Val Asp Asp Glu  
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 385 390 395 400  
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**SECRET**

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Gly Trp Ile Arg Glu Lys Asp Asn Asp Leu His Trp Glu Pro Ile Arg  
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Asp Pro Ala Gly Gly Gln Tyr Leu Thr Val Ser Ala Ala Lys Ala Pro  
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Gly Gly Lys Ala Ala Arg Leu Val Leu Pro Leu Gly Arg Leu Met His  
435 440 445

Ser Gly Asp Leu Cys Leu Ser Phe Arg His Lys Val Thr Gly Leu His  
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Ser Gly Thr Leu Gln Val Phe Val Arg Lys His Gly Ala His Gly Ala  
465 470 475 480

Ala Leu Trp Gly Arg Asn Gly Gly His Gly Trp Arg Gln Thr Gln Ile  
485 490 495

Thr Leu Arg Gly Ala Asp Ile Lys Ser Val Val Phe Lys Gly Glu Lys  
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Lys Gly His Cys Ser Glu Glu Arg  
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17





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Glu Ser Lys Gln Asn Leu Asn Val Asp Lys Asp Thr Thr Leu Pro Ala  
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 Ser Ala Arg Lys Val Lys Ser Ser Glu Ser Lys Ile Cys Val Leu Leu  
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 165 170 175  
 Ala Gln Leu Glu Glu Asn Leu Lys Glu Lys Asn Asp Glu Ile Leu Ser  
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 Leu Lys Gln Ser Leu Glu Asp Asn Ile Val Ile Leu Ser Lys Gln Val  
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 340 345 350



Ser Ala Ala His Thr Gln Ala Thr Leu Leu Leu Gln Glu Lys Tyr Asp  
355 360 365

Ser Met Val Gln Ser Leu Glu Asp Val Thr Ala Gln Phe Glu Ser Tyr  
370 375 380

Lys Ala Leu Thr Ala Ser Glu Ile Glu Asp Leu Lys Leu Glu Asn Ser  
385 390 395 400

Ser Leu Gln Glu Lys Ala Ala Lys Ala Gly Lys Asn Ala Glu Asp Val  
405 410 415

Gln His Gln Ile Leu Ala Thr Glu Ser Ser Asn Gln Glu Tyr Val Arg  
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Met Leu Leu Asp Leu Gln Thr Lys Ser Ala Leu Lys Glu Thr Glu Ile  
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Lys Glu Ile Thr Val Ser Phe Leu Gln Lys Ile Thr Asp Leu Gln Asn  
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Gln Leu Lys Gln Gln Glu Glu Asp Phe Arg Lys Gln Leu Glu Asp Glu  
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Glu Gly Arg Lys Ala Glu Lys Glu Asn Thr Thr Ala Glu Leu Thr Glu  
485 490 495

Glu Ile Asn Lys Trp Arg Leu Leu Tyr Glu Glu Leu Tyr Asn Lys Thr  
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Lys Pro Phe Gln Leu Gln Leu Asp Ala Phe Glu Val Glu Lys Gln Ala  
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Asp Ser Tyr Ala Lys Leu Leu Gly His Gln Asn Leu Lys Gln Lys Ile  
545 550 555 560

Lys His Val Val Lys Leu Lys Asp Glu Asn Ser Gln Leu Lys Ser Glu  
565 570 575

Val Ser Lys Leu Arg Cys Gln Leu Ala Lys Lys Lys Gln Ser Glu Thr  
580 585 590

Lys Leu Gln Glu Glu Leu Asn Lys Val Leu Gly Ile Lys His Phe Asp  
595 600 605

Pro Ser Lys Ala Phe His His Glu Ser Lys Glu Asn Phe Ala Leu Lys  
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Cys Gln Glu Ser Trp Lys  
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SECRET

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Ser	Leu	Ser	Ala	Asn	Asn	Ala	Thr	Leu	Glu	Lys	Gln	Leu	Ile	Glu	Leu					
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Thr	Arg	Thr	Asn	Glu	Leu	Leu	Lys	Ser	Lys	Phe	Ser	Glu	Asn	Asp	Asn					
	130					135					140									
Gln	Lys	Asn	Leu	Arg	Ile	Leu	Ser	Leu	Glu	Leu	Met	Lys	Leu	Arg	Asn					
	145				150					155					160					
Lys	Arg	Glu	Thr	Lys	Met	Arg	Gly	Met	Met	Ala	Lys	Gln	Glu	Gly	Met					
				165				170						175						
Glu	Met	Lys	Leu	Gln	Val	Thr	Gln	Arg	Ser	Leu	Glu	Glu	Ser	Gln	Gly					
			180					185					190							
Lys	Ile	Ala	Gln	Leu	Glu	Gly	Lys	Leu	Val	Ser	Ile	Glu	Lys	Glu	Lys					
		195						200				205								
Ile	Asp	Glu	Lys	Ser	Glu	Thr	Glu	Lys	Leu	Leu	Glu	Tyr	Ile	Glu	Glu					
	210					215					220									
Ile	Ser	Cys	Ala	Ser	Asp	Gln	Val	Glu	Lys	Tyr	Lys	Leu	Asp	Ile	Ala					
	225				230					235					240					
Gln	Leu	Glu	Glu	Asn	Leu	Lys	Glu	Lys	Asn	Asp	Glu	Ile	Leu	Ser	Leu					
				245					250					255						
Lys	Gln	Ser	Leu	Glu	Glu	Asn	Ile	Val	Ile	Leu	Ser	Lys	Gln	Val	Glu					
			260					265						270						
Asp	Leu	Asn	Val	Lys	Cys	Gln	Leu	Leu	Glu	Lys	Glu	Lys	Glu	Asp	His					
		275						280				285								
Val	Asn	Arg	Asn	Arg	Glu	His	Asn	Glu	Asn	Leu	Asn	Ala	Glu	Met	Gln					
	290					295					300									
Asn	Leu	Lys	Gln	Lys	Phe	Ile	Leu	Glu	Gln	Gln	Glu	Arg	Glu	Lys	Leu					
	305				310					315					320					
Gln	Gln	Lys	Glu	Leu	Gln	Ile	Asp	Ser	Leu	Leu	Gln	Gln	Glu	Lys	Glu					

325										330					335				
Leu	Ser	Ser	Ser	Leu	His	Gln	Lys	Leu	Cys	Ser	Phe	Gln	Glu	Glu	Met				
			340					345					350						
Val	Lys	Glu	Lys	Asn	Leu	Phe	Glu	Glu	Glu	Leu	Lys	Gln	Thr	Leu	Asp				
		355					360					365							
Glu	Leu	Asp	Lys	Leu	Gln	Gln	Lys	Glu	Glu	Gln	Ala	Glu	Arg	Leu	Val				
	370					375					380								
Lys	Gln	Leu	Glu	Glu	Glu	Ala	Lys	Ser	Arg	Ala	Glu	Glu	Leu	Lys	Leu				
385					390					395					400				
Leu	Glu	Glu	Lys	Leu	Lys	Gly	Lys	Glu	Ala	Glu	Leu	Glu	Lys	Ser	Ser				
				405					410					415					
Ala	Ala	His	Thr	Gln	Ala	Thr	Leu	Leu	Leu	Gln	Glu	Lys	Tyr	Asp	Ser				
			420					425						430					
Met	Val	Gln	Ser	Leu	Glu	Asp	Val	Thr	Ala	Gln	Phe	Glu	Gly	Tyr	Lys				
		435					440					445							
Ala	Leu	Thr	Ala	Ser	Glu	Ile	Glu	Asp	Leu	Lys	Leu	Glu	Asn	Ser	Ser				
	450					455					460								
Leu	Gln	Glu	Lys	Ala	Ala	Lys	Ala	Gly	Lys	Asn	Ala	Glu	Asp	Val	Gln				
465					470					475					480				
His	Gln	Ile	Leu	Ala	Thr	Glu	Ser	Ser	Asn	Gln	Glu	Tyr	Val	Arg	Met				
				485					490					495					
Leu	Leu	Asp	Leu	Gln	Thr	Lys	Ser	Ala	Leu	Lys	Glu	Thr	Glu	Ile	Lys				
			500					505					510						
Glu	Ile	Thr	Val	Ser	Phe	Leu	Gln	Lys	Ile	Thr	Asp	Leu	Gln	Asn	Gln				
		515					520					525							
Leu	Lys	Gln	Gln	Glu	Glu	Asp	Phe	Arg	Lys	Gln	Leu	Glu	Asp	Glu	Glu				
	530					535					540								
Gly	Arg	Lys	Ala	Glu	Lys	Glu	Asn	Thr	Thr	Ala	Glu	Leu	Thr	Glu	Glu				
545					550					555					560				
Ile	Asn	Lys	Trp	Arg	Leu	Leu	Tyr	Glu	Glu	Leu	Tyr	Asn	Lys	Thr	Lys				
				565					570					575					
Pro	Phe	Gln	Leu	Gln	Leu	Asp	Ala	Phe	Glu	Val	Glu	Lys	Gln	Ala	Leu				
			580					585						590					
Leu	Asn	Glu	His	Gly	Ala	Ala	Gln	Glu	Gln	Leu	Asn	Lys	Ile	Arg	Asp				
		595					600					605							
Ser	Tyr	Ala	Lys	Leu	Leu	Gly	His	Gln	Asn	Leu	Lys	Gln	Lys	Ile	Lys				
	610					615					620								
His	Val	Val	Lys	Leu	Lys	Asp	Glu	Asn	Ser	Gln	Leu	Lys	Ser	Glu	Val				





Glu Phe Asn Ala Asn His Pro Phe Leu Phe Phe Ile Arg His Asn Lys  
340 345 350

Thr Gln Thr Ile Leu Phe Tyr Gly Arg Val Cys Ser Pro  
355 360 365

<210> 21  
<211> 1590  
<212> DNA  
<213> Homo sapiens

<400> 21  
cagctcccgg gcaccatgcg aaccgccccg agcctccgcc gctgcgtctg cctgctgctc 60  
gccgcgatcc tggacctggc gcgctacctg acagtcaaca ttgagcctct ccccccctgtg 120  
gtggctggag acgccgtgac tttgaagtgt aacttcaaga cagatgggag catgaggag 180  
atcgtgtggt accgggtgac ggatgggtgc accatcaagc aaaagatctt caccttcgac 240  
gccatgttct ccaccaacta ctcacacatg gagaactacc gcaagcgaga ggacctgggtg 300  
taccagtcca ctgtgaggct gcccgaggct cggatctcag acaatggtcc ctatgagtgc 360  
catgtgggca tctacgaccg cgccaccagg gagaagggtg tcctggcatc aggcaacatc 420  
ttcctcaacg tcatggctcc tcccacctcc attgaagtgg tggctgctga cacaccagcc 480  
cccttcagcc gctaccaagc ccagaacttc acgctgggtc gcacgtgtgc tggaggaaaa 540  
ccagcaccca tggtttattt caaacgagat ggggaaccaa tcgacgcagt gcccttatca 600  
gagccaccag ctgcgagctc cggcccccta caggacagca ggcccttcg cagccttctg 660  
ctggacctgg atgacaccaa gatgcagaag tcactgtccc tcctggacgc cgagaaccgg 720  
ggtagggcgac cctacacgga gcgccccctc cgtggcctga ccccgatcc caacatcctc 780  
ctccagccaa ccacagagaa cataccagag acggtcgtga gccgtgagtt tccccgctgg 840  
gtccacagcg ccgagccccc ctacttctct cgccacagcc gcaccccgag cagtgcgggc 900  
actgtggaag tacgtgccct gctcacctgg accctcaacc cacagatcga caacgaggcc 960  
ctcttcagct gcgaggtcaa gcacccagct ctgtcgatgc ccatgcgggc agaggtcacg 1020  
ccggttgccc ccaaaggacc caaaattgtg atgacgccc gcagagcccc ggtaggggac 1080  
acagtgagga ttctgggtcca tgggtttcag aacgaagtct tcccggagcc catgttcacg 1140  
tggacgcggg ttgggagccg cctcctggac ggcagcgctg agttcgacgg gaaggagctg 1200  
gtgctggagc ggggttccgc cgagctcaat ggctccatgt atcgtctgcac cgcccagaa 1260  
ccactgggct ccaccgacac gcacaccgag ctcatcgtgt ttgaaaaccc aaatatccca 1320  
agaggaacgg aggaactctaa tgggttcatt ggccccactg gtgcccggct caccttggtg 1380  
ctcgccctga cagtgattct ggagctgacg tgaaggcacc cgccccggcc actccatcag 1440  
gcactgacat ctccacgacc ggttttcatt tcttttctaa actatttcca gtcttgttct 1500  
tagtctcttt ccactctgtg cttggcttct tcagtcgggt taattaaaac aaacagaaca 1560  
attttcccca aaaaaaaaaa aaaaaaaaaa 1590

<210> 22  
<211> 465  
<212> PRT  
<213> Homo sapiens

<400> 22  
Met Arg Thr Ala Pro Ser Leu Arg Arg Cys Val Cys Leu Leu Leu Ala  
1 5 10 15  
Ala Ile Leu Asp Leu Ala Arg Tyr Leu Thr Val Asn Ile Glu Pro Leu  
20 25 30  
Pro Pro Val Val Ala Gly Asp Ala Val Thr Leu Lys Cys Asn Phe Lys  
35 40 45

Thr Asp Gly Arg Met Arg Glu Ile Val Trp Tyr Arg Val Thr Asp Gly  
50 55 60

Gly Thr Ile Lys Gln Lys Ile Phe Thr Phe Asp Ala Met Phe Ser Thr  
65 70 75 80

Asn Tyr Ser His Met Glu Asn Tyr Arg Lys Arg Glu Asp Leu Val Tyr  
85 90 95

Gln Ser Thr Val Arg Leu Pro Glu Val Arg Ile Ser Asp Asn Gly Pro  
100 105 110

Tyr Glu Cys His Val Gly Ile Tyr Asp Arg Ala Thr Arg Glu Lys Val  
115 120 125

Val Leu Ala Ser Gly Asn Ile Phe Leu Asn Val Met Ala Pro Pro Thr  
130 135 140

Ser Ile Glu Val Val Ala Ala Asp Thr Pro Ala Pro Phe Ser Arg Tyr  
145 150 155 160

Gln Ala Gln Asn Phe Thr Leu Val Cys Ile Val Ser Gly Gly Lys Pro  
165 170 175

Ala Pro Met Val Tyr Phe Lys Arg Asp Gly Glu Pro Ile Asp Ala Val  
180 185 190

Pro Leu Ser Glu Pro Pro Ala Ala Ser Ser Gly Pro Leu Gln Asp Ser  
195 200 205

Arg Pro Phe Arg Ser Leu Leu Leu Asp Leu Asp Asp Thr Lys Met Gln  
210 215 220

Lys Ser Leu Ser Leu Leu Asp Ala Glu Asn Arg Gly Gly Arg Pro Tyr  
225 230 235 240

Thr Glu Arg Pro Ser Arg Gly Leu Thr Pro Asp Pro Asn Ile Leu Leu  
245 250 255

Gln Pro Thr Thr Glu Asn Ile Pro Glu Thr Val Val Ser Arg Glu Phe  
260 265 270

Pro Arg Trp Val His Ser Ala Glu Pro Thr Tyr Phe Leu Arg His Ser  
275 280 285

Arg Thr Pro Ser Ser Asp Gly Thr Val Glu Val Arg Ala Leu Leu Thr  
290 295 300

Trp Thr Leu Asn Pro Gln Ile Asp Asn Glu Ala Leu Phe Ser Cys Glu  
305 310 315 320

Val Lys His Pro Ala Leu Ser Met Pro Met Arg Ala Glu Val Thr Pro  
325 330 335

Val Ala Pro Lys Gly Pro Lys Ile Val Met Thr Pro Ser Arg Ala Arg  
340 345 350



Val Gly Asp Thr Val Arg Ile Leu Val His Gly Phe Gln Asn Glu Val  
355 360 365

Phe Pro Glu Pro Met Phe Thr Trp Thr Arg Val Gly Ser Arg Leu Leu  
370 375 380

Asp Gly Ser Ala Glu Phe Asp Gly Lys Glu Leu Val Leu Glu Arg Val  
385 390 395 400

Pro Ala Glu Leu Asn Gly Ser Met Tyr Arg Cys Thr Ala Gln Asn Pro  
405 410 415

Leu Gly Ser Thr Asp Thr His Thr Arg Leu Ile Val Phe Glu Asn Pro  
420 425 430

Asn Ile Pro Arg Gly Thr Glu Asp Ser Asn Gly Ser Ile Gly Pro Thr  
435 440 445

Gly Ala Arg Leu Thr Leu Val Leu Ala Leu Thr Val Ile Leu Glu Leu  
450 455 460

Thr  
465

<210> 23  
<211> 1593  
<212> DNA  
<213> Homo sapiens

<400> 23  
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gcccgcatcc tggacctggc gcgcggctac ctgacagtca acattgagcc tctccccct 120  
gtggtggctg gagacgccgt gactttgaag tgtaacttca agacagatgg gcgcatgcgg 180  
gagatcgtgt ggtaccgggt gacggatggt ggcaccatca agcaaaagat cttcaccttc 240  
gacgccatgt tctccaccaa ctactcacac atggagaact accgcaagcg agaggacctg 300  
gtgtaccagt ccactgtgag gctgccccgag gtccggatct cagacaatgg tccctatgag 360  
tgccatgtgg gcatctacga ccgcgccacc agggagaagg tggctcctggc atcaggcaac 420  
atcttctca acgtcatggc tctcccacc tccattgaag tgggtggctg tgacacacca 480  
gcccccttca gccgctacca agcccagaac ttcacgctgg tctgcatcgt gtctggagga 540  
aaaccagcac ccatggttta tttcaaacga gatggggaac caatcgacgc agtgccccta 600  
tcagagccac cagctgcgag ctccggcccc ctacaggaca gcaggccctt ccgcagcctt 660  
ctgctggacc tggatgacac caagatgcag aagtcactgt ccctcctgga cgccgagaa 720  
cggggtgggc gaccctacac ggagcgcccc tcccgtggcc tgaccccaga tcccaacatc 780  
ctcctccagc caaccacaga gaacatacca gagacggctg tgagccgtga gtttccccgc 840  
tgggtccaca gcgccgagcc cacctaactc ctgcgccaca gccgcacccc gagcagtga 900  
ggcactgtgg aagtacgtgc cctgctcacc tggaccctca acccacagat cgacaacgag 960  
gccctcttca gctgcgaggt caagcaccca gctctgtcga tgcccatgcg ggcagagggtc 1020  
acgcgggttg ccccaaaagg acccaaaatt gtgatgacgc ccagcagagc ccgggtagg 1080  
gacacagtga ggattctggt ccatgggttt cagaacgaag tcttcccga gcccatgttc 1140  
acgtggacgc ggggtgggag ccgcctcctg gacggcagcg ctgagttcga cgggaaggag 1200  
ctggtgctgg agcgggttcc cgccgagctc aatggctcca tgtatcgctg caccgcccag 1260  
aaccactgg gctccaccga cagcacacc cggctcatcg tgtttgaaaa cccaaatatc 1320  
ccaagaggaa cggaggactc taatggttcc attggccca ctggtgccc gctcacttg 1380  
gtgctcgccc tgacagtgat tctggagctg acgtgaaggc acccgccccg gccactccat 1440  
caggcactga catctccacg accggttttc atttcttttc taaactatct ccagtcttgt 1500

tcttagtctc ttccatctg tgtcttggt tcttcagtcg gtttaattaa aacaaacaga 1560  
acaattttcc ccaaaaaaaaa aaaaaaaaaa aaa 1593

<210> 24  
<211> 466  
<212> PRT  
<213> Homo sapiens

<400> 24  
Met Arg Thr Ala Pro Ser Leu Arg Arg Cys Val Cys Leu Leu Leu Ala  
1 5 10 15  
Ala Ile Leu Asp Leu Ala Arg Gly Tyr Leu Thr Val Asn Ile Glu Pro  
20 25 30  
Leu Pro Pro Val Val Ala Gly Asp Ala Val Thr Leu Lys Cys Asn Phe  
35 40 45  
Lys Thr Asp Gly Arg Met Arg Glu Ile Val Trp Tyr Arg Val Thr Asp  
50 55 60  
Gly Gly Thr Ile Lys Gln Lys Ile Phe Thr Phe Asp Ala Met Phe Ser  
65 70 75 80  
Thr Asn Tyr Ser His Met Glu Asn Tyr Arg Lys Arg Glu Asp Leu Val  
85 90 95  
Tyr Gln Ser Thr Val Arg Leu Pro Glu Val Arg Ile Ser Asp Asn Gly  
100 105 110  
Pro Tyr Glu Cys His Val Gly Ile Tyr Asp Arg Ala Thr Arg Glu Lys  
115 120 125  
Val Val Leu Ala Ser Gly Asn Ile Phe Leu Asn Val Met Ala Pro Pro  
130 135 140  
Thr Ser Ile Glu Val Val Ala Ala Asp Thr Pro Ala Pro Phe Ser Arg  
145 150 155 160  
Tyr Gln Ala Gln Asn Phe Thr Leu Val Cys Ile Val Ser Gly Gly Lys  
165 170 175  
Pro Ala Pro Met Val Tyr Phe Lys Arg Asp Gly Glu Pro Ile Asp Ala  
180 185 190  
Val Pro Leu Ser Glu Pro Pro Ala Ala Ser Ser Gly Pro Leu Gln Asp  
195 200 205  
Ser Arg Pro Phe Arg Ser Leu Leu Leu Asp Leu Asp Asp Thr Lys Met  
210 215 220  
Gln Lys Ser Leu Ser Leu Leu Asp Ala Glu Asn Arg Gly Gly Arg Pro  
225 230 235 240  
Tyr Thr Glu Arg Pro Ser Arg Gly Leu Thr Pro Asp Pro Asn Ile Leu  
245 250 255

Leu Gln Pro Thr Thr Glu Asn Ile Pro Glu Thr Val Val Ser Arg Glu  
260 265 270

Phe Pro Arg Trp Val His Ser Ala Glu Pro Thr Tyr Phe Leu Arg His  
275 280 285

Ser Arg Thr Pro Ser Ser Asp Gly Thr Val Glu Val Arg Ala Leu Leu  
290 295 300

Thr Trp Thr Leu Asn Pro Gln Ile Asp Asn Glu Ala Leu Phe Ser Cys  
305 310 315 320

Glu Val Lys His Pro Ala Leu Ser Met Pro Met Arg Ala Glu Val Thr  
325 330 335

Pro Val Ala Pro Lys Gly Pro Lys Ile Val Met Thr Pro Ser Arg Ala  
340 345 350

Arg Val Gly Asp Thr Val Arg Ile Leu Val His Gly Phe Gln Asn Glu  
355 360 365

Val Phe Pro Glu Pro Met Phe Thr Trp Thr Arg Val Gly Ser Arg Leu  
370 375 380

Leu Asp Gly Ser Ala Glu Phe Asp Gly Lys Glu Leu Val Leu Glu Arg  
385 390 395 400

Val Pro Ala Glu Leu Asn Gly Ser Met Tyr Arg Cys Thr Ala Gln Asn  
405 410 415

Pro Leu Gly Ser Thr Asp Thr His Thr Arg Leu Ile Val Phe Glu Asn  
420 425 430

Pro Asn Ile Pro Arg Gly Thr Glu Asp Ser Asn Gly Ser Ile Gly Pro  
435 440 445

Thr Gly Ala Arg Leu Thr Leu Val Leu Ala Leu Thr Val Ile Leu Glu  
450 455 460

Leu Thr  
465

<210> 25  
<211> 1407  
<212> DNA  
<213> Homo sapiens

<400> 25  
atgcgaaccg ccccgagcct ccgccgctgc gtctgcctgc tgctcgccgc gatcctggac 60  
ctggcgcgcg gctacctgac agtcaacatt gagcctctcc cccctgtggt ggctggagac 120  
gccgtgactt tgaagtgtaa cttcaagaca gatgggcgca tgcgggagat cgtgtggtac 180  
cgggtgacgg atggtggcac catcaagcaa aagatcttca ccttcgacgc catgttctcc 240  
accaactact cacacatgga gaactaccgc aagcgagagg acctggtgta ccagtccact 300  
gtgaggctgc ccgaggtccg gatctcagac aatggtccct atgagtgcca tgtgggcatc 360  
tacgaccgcg ccaccaggga gaaggtggtc ctggcatcag gcaacatctt cctcaacgtc 420

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atggctcctc ccacctccat tgaagtgggtg gctgctgaca caccagcccc cttcagccgc 480
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gtttattttca aacgagatgg ggaaccaatc gacgcagtgc ccctatcaga gccaccagct 600
gcgagctccg gccccctaca ggacagcagg cccttccgca gccttctgca ccgtgacctg 660
gatgacacca agatgcagaa gtcactgtcc ctccctggacg ccgagaaccg ggggtggcga 720
ccctacacgg agcgcctcct ccgtggcctg accccagatc ccaacatcct cctccagcca 780
accacagaga acataaccaga gacggctcgtg agccgtgagt ttccccgctg ggtccacagc 840
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cgggttcccg ccgagctcaa tggctccatg tatcgctgca ccgcccagaa cccactgggc 1260
tccaccgaca cgcacaccgg gctcatcgtg tttgaaaacc caaatatccc aagaggaacg 1320
gaggactcta atggttccat tggccccact ggtgcccggc tcaccttggg gctcgccctg 1380
acagtgatcc tggagctgac gtgaagg 1407

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<210> 26  
 <211> 467  
 <212> PRT  
 <213> Homo sapiens

<400> 26  
 Met Arg Thr Ala Pro Ser Leu Arg Arg Cys Val Cys Leu Leu Leu Ala  
 1 5 10 15  
 Ala Ile Leu Asp Leu Ala Arg Gly Tyr Leu Thr Val Asn Ile Glu Pro  
 20 25 30  
 Leu Pro Pro Val Val Ala Gly Asp Ala Val Thr Leu Lys Cys Asn Phe  
 35 40 45  
 Lys Thr Asp Gly Arg Met Arg Glu Ile Val Trp Tyr Arg Val Thr Asp  
 50 55 60  
 Gly Gly Thr Ile Lys Gln Lys Ile Phe Thr Phe Asp Ala Met Phe Ser  
 65 70 75 80  
 Thr Asn Tyr Ser His Met Glu Asn Tyr Arg Lys Arg Glu Asp Leu Val  
 85 90 95  
 Tyr Gln Ser Thr Val Arg Leu Pro Glu Val Arg Ile Ser Asp Asn Gly  
 100 105 110  
 Pro Tyr Glu Cys His Val Gly Ile Tyr Asp Arg Ala Thr Arg Glu Lys  
 115 120 125  
 Val Val Leu Ala Ser Gly Asn Ile Phe Leu Asn Val Met Ala Pro Pro  
 130 135 140  
 Thr Ser Ile Glu Val Val Ala Ala Asp Thr Pro Ala Pro Phe Ser Arg  
 145 150 155 160  
 Tyr Gln Ala Gln Asn Phe Thr Leu Val Cys Ile Val Ser Gly Gly Lys  
 165 170 175



<210> 27  
 <211> 682  
 <212> DNA  
 <213> Homo sapiens

<400> 27  
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 cgccacagcc gcaccccag cagtgcggc actgtggaag tacgtgccct gctcacctgg 180  
 accctcaacc cacagatcga caacgaggcc ctcttcagct gcgaggtcaa gcacccagct 240  
 ctgtcgatgc ccatgcgggc agaggtcacg ctggttgccc ccaaaggacc caaaattgtg 300  
 atgatgcccc gcagagcccg ggtaggggac acagtgcgga ttctggtcca tgggtttcag 360  
 aacgaagtct tcccggagcc catgttcacg tggacgcggg ttgggagccg cctcctggac 420  
 ggcagcgctg agttcgacgg gaaggagctg gtgctggagc gggttcccg cagactcaat 480  
 ggctccatgt atcgtgcac cgcccagaac ccactgggct ccactgacac gcacacccgg 540  
 ctcacgtgt ttgaaaaccc aaatatccca agaggaacgg aggactctaa tgggtccatt 600  
 gccccactg gtgcccggct caccttggtg ctgcacctga cagtgcattt ggagctgacg 660  
 tgatgacagt gattctggag ct 682

<210> 28  
 <211> 219  
 <212> PRT  
 <213> Homo sapiens

<400> 28  
 Met Arg Thr Ala Pro Ser Leu Arg Arg Cys Pro Pro Pro Pro Arg Pro  
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 Val Ala Gly Gly Arg Ser Ala Ser Glu Phe Pro Arg Trp Val His Ser  
 20 25 30  
 Ala Glu Pro Thr Tyr Phe Leu Arg His Ser Arg Thr Pro Ser Ser Asp  
 35 40 45  
 Gly Thr Val Glu Val Arg Ala Leu Leu Thr Trp Thr Leu Asn Pro Gln  
 50 55 60  
 Ile Asp Asn Glu Ala Leu Phe Ser Cys Glu Val Lys His Pro Ala Leu  
 65 70 75 80  
 Ser Met Pro Met Arg Ala Glu Val Thr Leu Val Ala Pro Lys Gly Pro  
 85 90 95  
 Lys Ile Val Met Met Pro Ser Arg Ala Arg Val Gly Asp Thr Val Arg  
 100 105 110  
 Ile Leu Val His Gly Phe Gln Asn Glu Val Phe Pro Glu Pro Met Phe  
 115 120 125  
 Thr Trp Thr Arg Val Gly Ser Arg Leu Leu Asp Gly Ser Ala Glu Phe  
 130 135 140  
 Asp Gly Lys Glu Leu Val Leu Glu Arg Val Pro Ala Glu Leu Asn Gly  
 145 150 155 160

Ser Met Tyr Arg Cys Thr Ala Gln Asn Pro Leu Gly Ser Thr Asp Thr  
165 170 175

His Thr Arg Leu Ile Val Phe Glu Asn Pro Asn Ile Pro Arg Gly Thr  
180 185 190

Glu Asp Ser Asn Gly Ser Ile Ala Pro Thr Gly Ala Arg Leu Thr Leu  
195 200 205

Val Leu Ala Leu Thr Val Ile Leu Glu Leu Thr  
210 215

<210> 29  
<211> 992  
<212> DNA  
<213> Homo sapiens

<400> 29  
accatgcgaa ccgccccgag cctccgcccgc tgcgtctgcc tgctgctcgc cgcgatacctg 60  
gacctggcgc gcggctacct gacagtcaac attgagcctc tccccctgt ggtggctgga 120  
gacgccgtga ctttgaagtg taacttcaag acagatgggc gcatgcggga gatcgtgtgg 180  
taccgggtga cggatgggtg caccatcaag caaaagatct tcacctcga cgccatgttc 240  
tccaccaact actcacacat ggagaactac cgcaagcgag aggacctggt gtaccagtcc 300  
actgtgaggc tgcccagagt ccgcatctca gacaatggtc cctatgagt ccattgtggc 360  
atctacgacc gcgccaccag ggagaagggt gtcctggcat caggcaacat cttcctcaac 420  
gtcatggttg ccccaaaagg acccaaaatt gtgatgacgc ccagcagagc ccgggtaggg 480  
gacacagtga ggattctggt ccatgggttt cagaacgaag tcttcccga gcccatgttc 540  
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ctgggtgctgg agcgggttcc cgccgagctc aatggctcca tgtatcgctg caccgccccg 660  
aaccactggt gctccaccga cagcacacc cggtcatcg tgtttgaaaa cccaaatattc 720  
ccaagaggaa cggaggactc taatgggttc attggcccca ctggtgccc gctcaccttg 780  
gtgctcgccc tgacagtgat tctggagctg acgtgaagac agtgattctg gactgacgtg 840  
gacagtgatt ctggagctga cgtgatgaca gtgattctgg agctgacgtg atgacagtga 900  
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<211> 270  
<212> PRT  
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<400> 30  
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1 5 10 15

Ala Ile Leu Asp Leu Ala Arg Gly Tyr Leu Thr Val Asn Ile Glu Pro  
20 25 30

Leu Pro Pro Val Val Ala Gly Asp Ala Val Thr Leu Lys Cys Asn Phe  
35 40 45

Lys Thr Asp Gly Arg Met Arg Glu Ile Val Trp Tyr Arg Val Thr Asp  
50 55 60

Gly Gly Thr Ile Lys Gln Lys Ile Phe Thr Phe Asp Ala Met Phe Ser  
 65 70 75 80  
 Thr Asn Tyr Ser His Met Glu Asn Tyr Arg Lys Arg Glu Asp Leu Val  
 85 90 95  
 Tyr Gln Ser Thr Val Arg Leu Pro Glu Val Arg Ile Ser Asp Asn Gly  
 100 105 110  
 Pro Tyr Glu Cys His Val Gly Ile Tyr Asp Arg Ala Thr Arg Glu Lys  
 115 120 125  
 Val Val Leu Ala Ser Gly Asn Ile Phe Leu Asn Val Met Val Ala Pro  
 130 135 140  
 Lys Gly Pro Lys Ile Val Met Thr Pro Ser Arg Ala Arg Val Gly Asp  
 145 150 155 160  
 Thr Val Arg Ile Leu Val His Gly Phe Gln Asn Glu Val Phe Pro Glu  
 165 170 175  
 Pro Met Phe Thr Trp Thr Arg Val Gly Ser Arg Leu Leu Asp Gly Ser  
 180 185 190  
 Ala Glu Phe Asp Gly Lys Glu Leu Val Leu Glu Arg Val Pro Ala Glu  
 195 200 205  
 Leu Asn Gly Ser Met Tyr Arg Cys Thr Ala Pro Asn Pro Leu Gly Ser  
 210 215 220  
 Thr Asp Thr His Thr Arg Leu Ile Val Phe Glu Asn Pro Asn Ile Pro  
 225 230 235 240  
 Arg Gly Thr Glu Asp Ser Asn Gly Ser Ile Gly Pro Thr Gly Ala Arg  
 245 250 255  
 Leu Thr Leu Val Leu Ala Leu Thr Val Ile Leu Glu Leu Thr  
 260 265 270

<210> 31  
 <211> 1341  
 <212> DNA  
 <213> Homo sapiens

<400> 31  
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 acggatgggtg gcacatcaa gcaaaagatc ttcaccttcg acgccatgtt ctccaccaac 180  
 tactcacaca tggagaacta ccgcaagcga gaggacctgg tgtaccagtc cactgtgagg 240  
 ctgcccagagg tccggatctc agacaatggt ccctatgagt gccatgtggg catctacgac 300  
 cgcgccacca gggagaaggt ggtcctggca tcaggcaaca tcttcctcaa cgtcatggct 360  
 cctccacact ccattgaagt ggtggctgct gacacaccag ccccttcag ccgctaccaa 420  
 gcccagaact tcacgctggg ctgcatcgtg tctggaggaa aaccagcacc catggtttat 480  
 ttcaaacgag atggggaacc aatcgacgca gtgccctat cagagccacc agctgcgagc 540  
 tccggccccc tacaggacag caggcccttc cgcagccttc tgcaccgtga cctggatgac 600  
 accaagatgc agaagtcact gtccctcctg gacgccgaga accgggggtgg gcgaccctac 660



acggagcgcc cctcccgtgg cctgacccca gatcccaaca tcctcctcca gcccaaccaca 720  
gagaacatac cagagacggg cgtgagccgt gagtttcccc gctgggtcca cagcgccgag 780  
cccacctact tcctgcgcca cagecgcacc ccgagcagtg acggcactgt ggaagtacgt 840  
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cccgcgagc tcaatggctc catgtatcgc tgcaccgccc agaaccact gggctccacc 1200  
gacacgcaca cccggctcat cgtgtttgaa aacccaaata tccaagagg aacggaggac 1260  
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attctggagc tgacgctcga g 1341

<210> 32  
<211> 447  
<212> PRT  
<213> Homo sapiens

<400> 32  
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Gly Asp Ala Val Thr Leu Lys Cys Asn Phe Lys Thr Asp Gly Arg Met  
20 25 30  
Arg Glu Ile Val Trp Tyr Arg Val Thr Asp Gly Gly Thr Ile Lys Gln  
35 40 45  
Lys Ile Phe Thr Phe Asp Ala Met Phe Ser Thr Asn Tyr Ser His Met  
50 55 60  
Glu Asn Tyr Arg Lys Arg Glu Asp Leu Val Tyr Gln Ser Thr Val Arg  
65 70 75 80  
Leu Pro Glu Val Arg Ile Ser Asp Asn Gly Pro Tyr Glu Cys His Val  
85 90 95  
Gly Ile Tyr Asp Arg Ala Thr Arg Glu Lys Val Val Leu Ala Ser Gly  
100 105 110  
Asn Ile Phe Leu Asn Val Met Ala Pro Pro Thr Ser Ile Glu Val Val  
115 120 125  
Ala Ala Asp Thr Pro Ala Pro Phe Ser Arg Tyr Gln Ala Gln Asn Phe  
130 135 140  
Thr Leu Val Cys Ile Val Ser Gly Gly Lys Pro Ala Pro Met Val Tyr  
145 150 155 160  
Phe Lys Arg Asp Gly Glu Pro Ile Asp Ala Val Pro Leu Ser Glu Pro  
165 170 175  
Pro Ala Ala Ser Ser Gly Pro Leu Gln Asp Ser Arg Pro Phe Arg Ser  
180 185 190  
Leu Leu His Arg Asp Leu Asp Asp Thr Lys Met Gln Lys Ser Leu Ser

195	200	205
Leu Leu Asp Ala Glu Asn Arg Gly Gly Arg Pro Tyr Thr Glu Arg Pro 210 215 220		
Ser Arg Gly Leu Thr Pro Asp Pro Asn Ile Leu Leu Gln Pro Thr Thr 225 230 235 240		
Glu Asn Ile Pro Glu Thr Val Val Ser Arg Glu Phe Pro Arg Trp Val 245 250 255		
His Ser Ala Glu Pro Thr Tyr Phe Leu Arg His Ser Arg Thr Pro Ser 260 265 270		
Ser Asp Gly Thr Val Glu Val Arg Ala Leu Leu Thr Trp Thr Leu Asn 275 280 285		
Pro Gln Ile Asp Asn Glu Ala Leu Phe Ser Cys Glu Val Lys His Pro 290 295 300		
Ala Leu Ser Met Pro Met Gln Ala Glu Val Thr Leu Val Ala Pro Lys 305 310 315 320		
Gly Pro Lys Ile Val Met Thr Pro Ser Arg Ala Arg Val Gly Asp Thr 325 330 335		
Val Arg Ile Leu Val His Gly Phe Gln Asn Glu Val Phe Pro Glu Pro 340 345 350		
Met Phe Thr Trp Thr Arg Val Gly Ser Arg Leu Leu Asp Gly Ser Ala 355 360 365		
Glu Phe Asp Gly Lys Glu Leu Val Leu Glu Arg Val Pro Ala Glu Leu 370 375 380		
Asn Gly Ser Met Tyr Arg Cys Thr Ala Gln Asn Pro Leu Gly Ser Thr 385 390 395 400		
Asp Thr His Thr Arg Leu Ile Val Phe Glu Asn Pro Asn Ile Pro Arg 405 410 415		
Gly Thr Glu Asp Ser Asn Gly Ser Ile Gly Pro Thr Gly Ala Arg Leu 420 425 430		
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<210> 33  
 <211> 702  
 <212> DNA  
 <213> Homo sapiens

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 acggatgggtg gcaccatcaa gcaaaagatc ttcaccttgc acgcatgtt ctccaccaac 180





His Phe Met Glu Gln Arg Val Tyr Pro Ala Glu Pro Glu Leu Gln Ser  
 65 70 75 80  
 His Gln Ala Ser Ala Ala Arg Trp Ser Pro Ser Pro Leu Ile Glu Asp  
 85 90 95  
 Leu Lys Glu Lys Ala Lys Ala Glu Gly Leu Trp Asn Leu Phe Leu Pro  
 100 105 110  
 Leu Glu Ala Asp Pro Glu Lys Lys Tyr Gly Ala Gly Leu Thr Asn Val  
 115 120 125  
 Glu Tyr Ala His Leu Cys Glu Leu Met Gly Thr Ser Leu Tyr Ala Pro  
 130 135 140  
 Glu Val Cys Asn Cys Ser Ala Pro Asp Thr Gly Asn Met Glu Leu Leu  
 145 150 155 160  
 Val Arg Tyr Gly Thr Glu Ala Gln Lys Ala Arg Trp Leu Ile Pro Leu  
 165 170 175  
 Leu Glu Gly Lys Ala Arg Ser Cys Phe Ala Met Thr Glu Pro Gln Val  
 180 185 190  
 Ala Ser Ser Asp Ala Thr Asn Ile Glu Ala Ser Ile Arg Glu Glu Asp  
 195 200 205  
 Ser Phe Tyr Val Ile Asn Gly His Lys Trp Trp Ile Thr Gly Ile Leu  
 210 215 220  
 Asp Pro Arg Cys Gln Leu Cys Val Phe Met Gly Lys Thr Asp Pro His  
 225 230 235 240  
 Ala Pro Arg His Arg Gln Gln Ser Val Leu Leu Val Pro Met Asp Thr  
 245 250 255  
 Pro Gly Ile Lys Ile Ile Arg Pro Leu Thr Val Tyr Gly Leu Glu Asp  
 260 265 270  
 Ala Pro Gly Gly His Gly Glu Val Arg Phe Glu His Val Arg Val Pro  
 275 280 285  
 Lys Glu Asn Met Val Leu Gly Pro Gly Arg Gly Phe Glu Ile Ala Gln  
 290 295 300  
 Gly Arg Leu Gly Pro Gly Arg Ile His His Cys Met Arg Leu Ile Gly  
 305 310 315 320  
 Phe Ser Glu Arg Ala Leu Ala Leu Met Lys Ala Arg Val Lys Ser Arg  
 325 330 335  
 Leu Ala Phe Gly Lys Pro Leu Val Glu Gln Gly Thr Val Leu Ala Asp  
 340 345 350  
 Ile Ala Gln Ser Arg Val Glu Ile Glu Gln Ala Arg Leu Leu Val Leu  
 355 360 365



43







A musical score for the song 'The Rose Tree'. It features a single melodic line on a five-line staff. The key signature has one sharp (F#), indicating the key of D major. The time signature is 4/4. The melody is written in a simple, folk-like style with many eighth and sixteenth notes. The lyrics 'The Rose Tree' are written below the staff, aligned with the notes. The score is presented in a clean, black-and-white format.

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45





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gaacacagtc ctgtgcaaaa ttcacagtgt ggatctgcag gctgtgggac ttgaagacta 1200
tggaagcaa ggggactata ttccacgcca ggtacgaacc tgggttaagc agtatcgagc 1260
ttccgaaact agcaccatcc cagccatgga gaggtgatc gaatggctgc ccctccatct 1320
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cccccttgct gatgtggcct acagctgcct ggctcattac ctgccatcca gttttcccg 1500
gctgagaggt attaatact gtgacttgac acagctggga atccctgctg cagaggagta 1560
tttcaggatg tactgtctcc aaatggggct ccctccact gagaactgga acttctatat 1620
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gcacttcatg gagcaacgtg tgtaccctgc agagccagag ctgcagagtc accaggcctc 2040
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gggcagactg gggcccgcca ggatccatca ctgcatgagg ctgacgggt tctcagagag 2760
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aaaaaaaaa 3490

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 <211> 1080  
 <212> PRT  
 <213> Homo sapiens

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 Glu Val Gln Asn Arg Ile Pro Ser Gly Thr Ile Leu Lys Ala Leu Met  
           20                    25                    30  
 Glu Gly Gly Glu Asn Gly Pro Trp Met Arg Phe Met Arg Ala Glu Ile  
           35                    40                    45

Thr Ala Glu Gly Phe Leu Arg Glu Phe Gly Arg Leu Cys Ser Glu Met  
50 55 60

Leu Lys Thr Ser Val Pro Val Asp Ser Phe Phe Ser Leu Leu Thr Ser  
65 70 75 80

Glu Arg Val Ala Lys Gln Phe Pro Val Met Thr Glu Ala Ile Thr Gln  
85 90 95

Ile Arg Ala Lys Gly Leu Gln Thr Ala Val Leu Ser Asn Asn Phe Tyr  
100 105 110

Leu Pro Asn Gln Lys Ser Phe Leu Pro Leu Asp Arg Lys Gln Phe Asp  
115 120 125

Val Ile Val Glu Ser Cys Met Glu Gly Ile Cys Lys Pro Asp Pro Arg  
130 135 140

Ile Tyr Lys Leu Cys Leu Glu Gln Leu Gly Leu Gln Pro Ser Glu Ser  
145 150 155 160

Ile Phe Leu Asp Asp Leu Gly Thr Asn Leu Lys Glu Ala Ala Arg Leu  
165 170 175

Gly Ile His Thr Ile Lys Val Asn Asp Pro Glu Thr Ala Val Lys Glu  
180 185 190

Leu Glu Ala Leu Leu Gly Phe Thr Leu Arg Val Gly Val Pro Asn Thr  
195 200 205

Arg Pro Val Lys Lys Thr Met Glu Ile Pro Lys Asp Ser Leu Gln Lys  
210 215 220

Tyr Leu Lys Asp Leu Leu Gly Ile Gln Thr Thr Gly Pro Leu Glu Leu  
225 230 235 240

Leu Gln Phe Asp His Gly Gln Ser Asn Pro Thr Tyr Tyr Ile Arg Leu  
245 250 255

Ala Asn Arg Asp Leu Val Leu Arg Lys Lys Pro Pro Gly Thr Leu Leu  
260 265 270

Pro Ser Ala His Ala Ile Glu Arg Glu Phe Arg Ile Met Lys Ala Leu  
275 280 285

Ala Asn Ala Gly Val Pro Val Pro Asn Val Leu Asp Leu Cys Glu Asp  
290 295 300

Ser Ser Val Ile Gly Thr Pro Phe Tyr Val Met Glu Tyr Cys Pro Gly  
305 310 315 320

Leu Ile Tyr Lys Asp Pro Ser Leu Pro Gly Leu Glu Pro Ser His Arg  
325 330 335

Arg Ala Ile Tyr Thr Ala Met Asn Thr Val Leu Cys Lys Ile His Ser  
340 345 350

Val Asp Leu Gln Ala Val Gly Leu Glu Asp Tyr Gly Lys Gln Gly Asp  
355 360 365

Tyr Ile Pro Arg Gln Val Arg Thr Trp Val Lys Gln Tyr Arg Ala Ser  
370 375 380

Glu Thr Ser Thr Ile Pro Ala Met Glu Arg Leu Ile Glu Trp Leu Pro  
385 390 395 400

Leu His Leu Pro Arg Gln Gln Arg Thr Thr Val Val His Gly Asp Phe  
405 410 415

Arg Leu Asp Asn Leu Val Phe His Pro Glu Glu Pro Glu Val Leu Ala  
420 425 430

Val Leu Asp Trp Glu Leu Ser Thr Leu Gly Asp Pro Leu Ala Asp Val  
435 440 445

Ala Tyr Ser Cys Leu Ala His Tyr Leu Pro Ser Ser Phe Pro Val Leu  
450 455 460

Arg Gly Ile Asn Asp Cys Asp Leu Thr Gln Leu Gly Ile Pro Ala Ala  
465 470 475 480

Glu Glu Tyr Phe Arg Met Tyr Cys Leu Gln Met Gly Leu Pro Pro Thr  
485 490 495

Glu Asn Trp Asn Phe Tyr Met Ala Phe Ser Phe Phe Arg Val Ala Ala  
500 505 510

Ile Leu Gln Gly Val Tyr Lys Arg Ser Leu Thr Gly Gln Ala Ser Ser  
515 520 525

Thr Tyr Ala Glu Gln Thr Gly Lys Leu Thr Glu Phe Val Ser Asn Leu  
530 535 540

Ala Trp Asp Phe Ala Val Lys Glu Gly Phe Arg Val Phe Lys Glu Met  
545 550 555 560

Pro Phe Thr Asn Pro Leu Thr Arg Ser Tyr His Thr Trp Ala Arg Pro  
565 570 575

Gln Ser Gln Trp Cys Pro Ile Gly Ser Arg Ser Tyr Ser Ser Val Pro  
580 585 590

Glu Ala Ser Pro Ala His Thr Ser Arg Gly Gly Leu Val Ile Ser Pro  
595 600 605

Glu Ser Leu Ser Pro Pro Val Arg Glu Leu Tyr His Arg Leu Lys His  
610 615 620

Phe Met Glu Gln Arg Val Tyr Pro Ala Glu Pro Glu Leu Gln Ser His  
625 630 635 640

Gln Ala Ser Ala Ala Arg Trp Ser Pro Ser Pro Leu Ile Glu Asp Leu  
645 650 655

Lys Glu Lys Ala Lys Ala Glu Gly Leu Trp Asn Leu Phe Leu Pro Leu  
 660 665 670  
 Glu Ala Asp Pro Glu Lys Lys Tyr Gly Ala Gly Leu Thr Asn Val Glu  
 675 680 685  
 Tyr Ala His Leu Cys Glu Leu Met Gly Thr Ser Leu Tyr Ala Pro Glu  
 690 695 700  
 Val Cys Asn Cys Ser Ala Pro Asp Thr Gly Asn Met Glu Leu Leu Val  
 705 710 715 720  
 Arg Tyr Gly Thr Glu Ala Gln Lys Ala Arg Trp Leu Ile Pro Leu Leu  
 725 730 735  
 Glu Gly Lys Ala Arg Ser Cys Phe Ala Met Thr Glu Pro Gln Val Ala  
 740 745 750  
 Ser Ser Asp Ala Thr Asn Ile Glu Ala Ser Ile Arg Glu Glu Asp Ser  
 755 760 765  
 Phe Tyr Val Ile Asn Gly His Lys Trp Trp Ile Thr Gly Ile Leu Asp  
 770 775 780  
 Pro Arg Cys Gln Leu Cys Val Phe Met Gly Lys Thr Asp Pro His Ala  
 785 790 795 800  
 Pro Arg His Arg Gln Gln Ser Val Leu Leu Val Pro Met Asp Thr Pro  
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 Gly Ile Lys Ile Ile Arg Pro Leu Thr Val Tyr Gly Leu Glu Asp Ala  
 820 825 830  
 Pro Gly Gly His Gly Glu Val Arg Phe Glu His Val Arg Val Pro Lys  
 835 840 845  
 Glu Asn Met Val Leu Gly Pro Gly Arg Gly Phe Glu Ile Ala Gln Gly  
 850 855 860  
 Arg Leu Gly Pro Gly Arg Ile His His Cys Met Arg Leu Ile Gly Phe  
 865 870 875 880  
 Ser Glu Arg Ala Leu Ala Leu Met Lys Ala Arg Val Lys Ser Arg Leu  
 885 890 895  
 Ala Phe Gly Lys Pro Leu Val Glu Gln Gly Thr Val Leu Ala Asp Ile  
 900 905 910  
 Ala Gln Ser Arg Val Glu Ile Glu Gln Ala Arg Leu Leu Val Leu Arg  
 915 920 925  
 Ala Ala His Leu Met Asp Leu Ala Gly Asn Lys Ala Ala Ala Leu Asp  
 930 935 940  
 Ile Ala Met Ile Lys Met Val Ala Pro Ser Met Ala Ser Arg Val Ile  
 945 950 955 960

Asp Arg Ala Ile Gln Lys Thr Ser Leu Gln Glu Ala Trp Ser Leu Phe  
965 970 975

Gln Ala Arg Arg Arg Gly Phe Ala Glu Gly Gln Gly Gly Ser Gly Thr  
980 985 990

Glu Ser Gly Lys Leu Val Phe Arg Leu Ser Val Pro Gly Trp Ala Gly  
995 1000 1005

Thr Val Thr Ser Leu Gln Pro Phe Ser Pro Ser Leu Ser Ala Cys Gly  
1010 1015 1020

Asn Leu Asp Thr Phe Trp Glu Ala Ser Gln Gly Cys Gly Thr Cys Leu  
1025 1030 1035 1040

Leu Trp Gln Leu Gln Gly Ser Cys Leu Ala Ser Leu Val Ser Arg Gly  
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Glu Asn Gly Met Gln Pro Thr Leu  
1075 1080

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35 40 45

Thr Ala Gly Asp Lys Leu Gln Pro Ser Pro Pro Pro Leu Ser Pro Pro  
50 55 60

Pro Arg Ala Pro Pro Leu Ser Pro Gly Pro Gly Gly Cys Phe Glu Gly  
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Gly Ala Gly Asn Cys Ser Ser Arg Gly Gly Arg Ala Ser Asp His Pro  
85 90 95

Gly Gly Gly Arg Glu Phe Phe Phe Asp Arg His Pro Gly Val Phe Ala  
100 105 110

Tyr Val Leu Asn Tyr Tyr Arg Thr Gly Lys Leu His Cys Pro Ala Asp  
115 120 125

Val Cys Gly Pro Leu Phe Glu Glu Glu Leu Ala Phe Trp Gly Ile Asp  
130 135 140

Glu Thr Asp Val Glu Pro Cys Cys Trp Met Thr Tyr Arg Gln His Arg  
145 150 155 160

Asp Ala Glu Glu Ala Leu Asp Ile Phe Glu Thr Pro Asp Leu Ile Gly  
165 170 175

Gly Asp Pro Gly Asp Asp Glu Asp Leu Ala Ala Lys Arg Leu Gly Ile  
180 185 190

Glu Asp Ala Ala Gly Leu Gly Gly Pro Asp Gly Lys Ser Gly Arg Trp  
195 200 205

Arg Arg Leu Gln Pro Arg Met Trp Ala Leu Phe Glu Asp Pro Tyr Ser  
210 215 220

Ser Arg Ala Ala Arg Phe Ile Ala Phe Ala Ser Leu Phe Phe Ile Leu  
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Val Ser Ile Thr Thr Phe Cys Leu Glu Thr His Glu Ala Phe Asn Ile  
245 250 255

Val Lys Asn Lys Thr Glu Pro Val Ile Asn Gly Thr Ser Val Val Leu  
260 265 270

Gln Tyr Glu Ile Glu Thr Asp Pro Ala Leu Thr Tyr Val Glu Gly Val  
275 280 285

Cys Val Val Trp Phe Thr Phe Glu Phe Leu Val Arg Ile Val Phe Ser  
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Pro Asn Lys Leu Glu Phe Ile Lys Asn Leu Leu Asn Ile Ile Asp Phe  
305 310 315 320

Val Ala Ile Leu Pro Phe Tyr Leu Glu Val Gly Leu Ser Gly Leu Ser  
325 330 335

Ser Lys Ala Ala Lys Asp Val Leu Gly Phe Leu Arg Val Val Arg Phe  
340 345 350

Val Arg Ile Leu Arg Ile Phe Lys Leu Thr Arg His Phe Val Gly Leu  
355 360 365

Arg Val Leu Gly His Thr Leu Arg Ala Ser Thr Asn Glu Phe Leu Leu  
370 375 380

Leu Ile Ile Phe Leu Ala Leu Gly Val Leu Ile Phe Ala Thr Met Ile  
385 390 395 400

Tyr Tyr Ala Glu Arg Val Gly Ala Gln Pro Asn Asp Pro Ser Ala Ser  
405 410 415

Glu His Thr Gln Phe Lys Asn Ile Pro Ile Gly Phe Trp Trp Ala Val  
420 425 430

Val Thr Met Thr Thr Leu Gly Tyr Gly Asp Met Tyr Pro Gln Thr Trp  
435 440 445







370 375 380

Leu Ile Ile Phe Leu Ala Leu Gly Val Leu Ile Phe Ala Thr Met Ile  
385 390 395 400

Tyr Tyr Ala Glu Arg Val Gly Ala Gln Pro Asn Asp Pro Ser Ala Ser  
405 410 415

Glu His Thr Gln Phe Lys Asn Ile Pro Ile Gly Phe Trp Trp Ala Val  
420 425 430

Val Thr Met Thr Thr Leu Gly Tyr Gly Asp Met Tyr Pro Gln Thr Trp  
435 440 445

Ser Gly Met Leu Val Gly Ala Leu Cys Ala Leu Ala Gly Val Leu Thr  
450 455 460

Ile Ala Met Pro Val Pro Val Ile Val Asn Asn Phe Gly Met Tyr Tyr  
465 470 475 480

Ser Leu Ala Met Ala Lys Gln Lys Leu Pro Arg Lys Arg Lys Lys His  
485 490 495

Ile Pro Pro Ala Pro Leu Ala Ser Ser Pro Thr Phe Cys Lys Thr Glu  
500 505 510

Leu Asn Met Ala Cys Asn Ser Thr Gln Ser Asp Thr Cys Leu Gly Lys  
515 520 525

Glu Asn Arg Leu Leu Glu His Asn Arg Ser Val Leu Ser Gly Asp Asp  
530 535 540

Ser Thr Gly Ser Glu Pro Pro Leu Ser Pro Pro Glu Arg Leu Pro Ile  
545 550 555 560

Arg Arg Ser Ser Thr Arg Asp Lys Asn Arg Arg Gly Glu Thr Cys Phe  
565 570 575

Leu Leu Thr Thr Gly Asp Tyr Thr Cys Ala Ser Asp Gly Gly Ile Arg  
580 585 590

Lys Gly Tyr Glu Lys Ser Arg Ser Leu Asn Asn Ile Ala Gly Leu Ala  
595 600 605

Gly Asn Ala Leu Arg Leu Ser Pro Val Thr Ser Pro Tyr Asn Ser Pro  
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Cys Pro Leu Arg Arg Ser Arg Ser Pro Ile Pro Ser Ile Leu  
625 630 635

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<213> Homo sapiens  
  
<400> 45

Met Gly Lys Ile Glu Asn Asn Glu Arg Val Ile Leu Asn Val Gly Gly  
 1 5 10 15  
 Thr Arg His Glu Thr Tyr Arg Ser Thr Leu Lys Thr Leu Pro Gly Thr  
 20 25 30  
 Arg Leu Ala Leu Leu Ala Ser Ser Glu Pro Pro Gly Asp Cys Leu Thr  
 35 40 45  
 Thr Ala Gly Asp Lys Leu Gln Pro Ser Pro Pro Pro Leu Ser Pro Pro  
 50 55 60  
 Pro Arg Ala Pro Pro Leu Ser Pro Gly Pro Gly Gly Cys Phe Glu Gly  
 65 70 75 80  
 Gly Ala Gly Asn Cys Ser Ser Arg Gly Gly Arg Ala Ser Asp His Pro  
 85 90 95  
 Gly Gly Gly Arg Glu Phe Phe Phe Asp Arg His Pro Gly Val Phe Ala  
 100 105 110  
 Tyr Val Leu Asn Tyr Tyr Arg Thr Gly Lys Leu His Cys Pro Ala Asp  
 115 120 125  
 Val Cys Gly Pro Leu Phe Glu Glu Glu Leu Ala Phe Trp Gly Ile Asp  
 130 135 140  
 Glu Thr Asp Val Glu Pro Cys Cys Trp Met Thr Tyr Arg Gln His Arg  
 145 150 155 160  
 Asp Ala Glu Glu Ala Leu Asp Ile Phe Glu Thr Pro Asp Leu Ile Gly  
 165 170 175  
 Gly Asp Pro Gly Asp Asp Glu Asp Leu Ala Ala Lys Arg Leu Gly Ile  
 180 185 190  
 Glu Asp Ala Ala Gly Leu Gly Gly Pro Asp Gly Lys Ser Gly Arg Trp  
 195 200 205  
 Arg Arg Leu Gln Pro Arg Met Trp Ala Leu Phe Glu Asp Pro Tyr Ser  
 210 215 220  
 Ser Arg Ala Ala Arg Phe Ile Ala Phe Ala Ser Leu Phe Phe Ile Leu  
 225 230 235 240  
 Val Ser Ile Thr Thr Phe Cys Leu Glu Thr His Glu Ala Phe Asn Ile  
 245 250 255  
 Val Lys Asn Lys Thr Glu Pro Val Ile Asn Gly Thr Ser Val Val Leu  
 260 265 270  
 Gln Tyr Glu Ile Glu Thr Asp Pro Ala Leu Thr Tyr Val Glu Gly Val  
 275 280 285  
 Cys Val Val Trp Phe Thr Phe Glu Phe Leu Val Arg Ile Val Phe Ser  
 290 295 300

Pro Asn Lys Leu Glu Phe Ile Lys Asn Leu Leu Asn Ile Ile Asp Phe  
305 310 315 320

Val Ala Ile Leu Pro Phe Tyr Leu Glu Val Gly Leu Ser Gly Leu Ser  
325 330 335

Ser Lys Ala Ala Lys Asp Val Leu Gly Phe Leu Arg Val Val Arg Phe  
340 345 350

Val Arg Ile Leu Arg Ile Phe Lys Leu Thr Arg His Phe Val Gly Leu  
355 360 365

Arg Val Leu Gly His Thr Leu Arg Ala Ser Thr Asn Glu Phe Leu Leu  
370 375 380

Leu Ile Ile Phe Leu Ala Leu Gly Val Leu Ile Phe Ala Thr Met Ile  
385 390 395 400

Tyr Tyr Ala Glu Arg Val Gly Ala Gln Pro Asn Asp Pro Ser Ala Ser  
405 410 415

Glu His Thr Gln Phe Lys Asn Ile Pro Ile Gly Phe Trp Trp Ala Val  
420 425 430

Val Thr Met Thr Thr Leu Gly Tyr Gly Asp Met Tyr Pro Gln Thr Trp  
435 440 445

Ser Gly Met Leu Val Gly Ala Leu Cys Ala Leu Ala Gly Val Leu Thr  
450 455 460

Ile Ala Met Pro Val Pro Val Ile Val Asn Asn Phe Gly Met Tyr Tyr  
465 470 475 480

Ser Leu Ala Met Ala Lys Gln Lys Leu Pro Arg Lys Arg Lys Lys His  
485 490 495

Ile Pro Pro Ala Pro Gln Ala Ser Ser Pro Thr Phe Cys Lys Thr Glu  
500 505 510

Leu Asn Met Ala Cys Asn Ser Thr Gln Ser Asp Thr Cys Leu Gly Lys  
515 520 525

Asp Asn Arg Leu Leu Glu His Asn Arg Ser Val Leu Ser Gly Asp Asp  
530 535 540

Ser Thr Gly Ser Glu Pro Pro Leu Ser Pro Pro Glu Arg Leu Pro Ile  
545 550 555 560

Arg Arg Ser Ser Thr Arg Asp Lys Asn Arg Arg Gly Glu Thr Cys Phe  
565 570 575

Leu Leu Thr Thr Gly Asp Tyr Thr Cys Ala Ser Asp Gly Gly Ile Arg  
580 585 590

Lys Asp Asn Cys Lys Glu Val Val Ile Thr Gly Tyr Thr Gln Ala Glu  
595 600 605



Val Lys Asn Lys Thr Glu Pro Val Ile Asn Gly Thr Ser Ala Val Leu  
260 265 270

Gln Tyr Glu Ile Glu Thr Asp Pro Ala Leu Thr Tyr Val Glu Gly Val  
275 280 285

Cys Val Val Trp Phe Thr Phe Glu Phe Leu Val Arg Ile Val Phe Ser  
290 295 300

Pro Asn Lys Leu Glu Phe Ile Lys Asn Leu Leu Asn Ile Ile Asp Phe  
305 310 315 320

Val Ala Ile Leu Pro Phe Tyr Leu Glu Val Gly Leu Ser Gly Leu Ser  
325 330 335

Ser Lys Ala Ala Lys Asp Val Leu Gly Phe Leu Arg Val Val Arg Phe  
340 345 350

Val Arg Ile Leu Arg Ile Phe Lys Leu Thr Arg His Phe Val Gly Leu  
355 360 365

Arg Val Leu Gly His Thr Leu Arg Ala Ser Thr Asn Glu Phe Leu Leu  
370 375 380

Leu Ile Ile Phe Leu Ala Leu Gly Val Leu Ile Phe Ala Thr Met Ile  
385 390 395 400

Tyr Tyr Ala Glu Arg Val Gly Ala Gln Pro Asn Asp Pro Ser Ala Ser  
405 410 415

Glu His Thr Gln Phe Lys Asn Ile Pro Ile Gly Phe Trp Trp Ala Val  
420 425 430

Val Thr Met Thr Thr Leu Gly Tyr Gly Asp Met Tyr Pro Gln Thr Trp  
435 440 445

Ser Gly Met Leu Val Gly Ala Leu Cys Ala Leu Ala Gly Val Leu Thr  
450 455 460

Ile Ala Met Pro Val Pro Val Ile Val Asn Asn Phe Gly Met Tyr Tyr  
465 470 475 480

Ser Leu Ala Met Ala Lys Gln Lys Leu Pro Arg Lys Arg Lys Lys His  
485 490 495

Ile Pro Pro Ala Pro Leu Ala Ser Ser Pro Thr Phe Cys Lys Thr Glu  
500 505 510

Leu Asn Met Ala Cys Asn Ser Thr Gln Ser Asp Thr Cys Leu Gly Lys  
515 520 525

Glu Asn Arg Leu Leu Glu His Asn Arg Ser Val Leu Ser Gly Asp Asp  
530 535 540

Ser Thr Gly Ser Glu Pro Pro Leu Ser Pro Pro Glu Arg Leu Pro Ile  
545 550 555 560

Arg Arg Ser Ser Thr Arg Asp Lys Asn Arg Arg Gly Glu Thr Cys Phe  
565 570 575

Leu Leu Thr Thr Gly Asp Tyr Thr Cys Ala Ser Asp Gly Gly Ile Arg  
580 585 590

Lys Asp Asn Cys Lys Asp Val Val Ile Thr Gly Tyr Thr Gln Ala Glu  
595 600 605

Ala Arg Ser Leu Thr  
610

<210> 47

<211> 624

<212> PRT

<213> Rattus norvegicus

<400> 47

Met Ser Lys Ile Glu Asn Asn Glu Arg Val Ile Leu Asn Val Gly Gly  
1 5 10 15

Thr Arg His Glu Thr Tyr Arg Ser Thr Leu Lys Thr Leu Pro Gly Thr  
20 25 30

Arg Leu Ala Leu Leu Ala Ser Ser Glu Pro Gln Gly Asp Cys Leu Thr  
35 40 45

Ala Ala Gly Asp Lys Leu Gln Pro Leu Pro Pro Pro Leu Ser Pro Pro  
50 55 60

Pro Arg Pro Pro Pro Leu Ser Pro Val Pro Ser Gly Cys Phe Glu Gly  
65 70 75 80

Gly Ala Gly Asn Cys Ser Ser His Gly Gly Asn Gly Ser Asp His Pro  
85 90 95

Gly Gly Gly Arg Glu Phe Phe Phe Asp Arg His Pro Gly Val Phe Ala  
100 105 110

Tyr Val Leu Asn Tyr Tyr Arg Thr Gly Lys Leu His Cys Pro Ala Asp  
115 120 125

Val Cys Gly Pro Leu Phe Glu Glu Leu Ala Phe Trp Gly Ile Asp  
130 135 140

Glu Thr Asp Val Glu Pro Cys Cys Trp Met Thr Tyr Arg Gln His Arg  
145 150 155 160

Asp Ala Glu Glu Ala Leu Asp Ile Phe Glu Thr Pro Asp Leu Ile Gly  
165 170 175

Gly Asp Pro Gly Asp Asp Glu Asp Leu Gly Gly Lys Arg Leu Gly Ile  
180 185 190

Glu Asp Ala Ala Gly Leu Gly Gly Pro Asp Gly Lys Ser Gly Arg Trp



195					200					205					
Arg	Lys	Leu	Gln	Pro	Arg	Met	Trp	Ala	Leu	Phe	Glu	Asp	Pro	Tyr	Ser
210					215					220					
Ser	Arg	Ala	Ala	Arg	Phe	Ile	Ala	Phe	Ala	Ser	Leu	Phe	Phe	Ile	Leu
225					230					235					240
Val	Ser	Ile	Thr	Thr	Phe	Cys	Leu	Glu	Thr	His	Glu	Ala	Phe	Asn	Ile
				245					250					255	
Val	Lys	Asn	Lys	Thr	Glu	Pro	Val	Ile	Asn	Gly	Thr	Ser	Ala	Val	Leu
			260					265					270		
Gln	Tyr	Glu	Ile	Glu	Thr	Asp	Pro	Ala	Leu	Thr	Tyr	Val	Glu	Gly	Val
		275					280					285			
Cys	Val	Val	Trp	Phe	Thr	Phe	Glu	Phe	Leu	Val	Arg	Ile	Val	Phe	Ser
290					295					300					
Pro	Asn	Lys	Leu	Glu	Phe	Ile	Lys	Asn	Leu	Leu	Asn	Ile	Ile	Asp	Phe
305					310					315					320
Val	Ala	Ile	Leu	Pro	Phe	Tyr	Leu	Glu	Val	Gly	Leu	Ser	Gly	Leu	Ser
				325					330					335	
Ser	Lys	Ala	Ala	Lys	Asp	Val	Leu	Gly	Phe	Leu	Arg	Val	Val	Arg	Phe
			340					345					350		
Val	Arg	Ile	Leu	Arg	Ile	Phe	Lys	Leu	Thr	Arg	His	Phe	Val	Gly	Leu
		355					360					365			
Arg	Val	Leu	Gly	His	Thr	Leu	Arg	Ala	Ser	Thr	Asn	Glu	Phe	Leu	Leu
		370					375					380			
Leu	Ile	Ile	Phe	Leu	Ala	Leu	Gly	Val	Leu	Ile	Phe	Ala	Thr	Met	Ile
385				390						395					400
Tyr	Tyr	Ala	Glu	Arg	Val	Gly	Ala	Gln	Pro	Asn	Asp	Pro	Ser	Ala	Ser
				405					410					415	
Glu	His	Thr	Gln	Phe	Lys	Asn	Ile	Pro	Ile	Gly	Phe	Trp	Trp	Ala	Val
			420					425					430		
Val	Thr	Met	Thr	Thr	Leu	Gly	Tyr	Gly	Asp	Met	Tyr	Pro	Gln	Thr	Trp
		435					440					445			
Ser	Gly	Met	Leu	Val	Gly	Ala	Leu	Cys	Ala	Leu	Ala	Gly	Val	Leu	Thr
		450				455					460				
Ile	Ala	Met	Pro	Val	Pro	Val	Ile	Val	Asn	Asn	Phe	Gly	Met	Tyr	Tyr
465				470					475					480	
Ser	Leu	Ala	Met	Ala	Lys	Gln	Lys	Leu	Pro	Arg	Lys	Arg	Lys	Lys	His
				485					490					495	
Ile	Pro	Pro	Ala	Pro	Leu	Ala	Ser	Ser	Pro	Thr	Phe	Cys	Lys	Thr	Glu

500					505					510					
Leu	Asn	Met	Ala	Cys	Asn	Ser	Thr	Gln	Ser	Asp	Thr	Cys	Leu	Gly	Lys
	515						520					525			
Glu	Asn	Arg	Leu	Leu	Glu	His	Asn	Arg	Ser	Val	Leu	Ser	Gly	Asp	Asp
	530					535					540				
Ser	Thr	Gly	Ser	Glu	Pro	Pro	Leu	Ser	Pro	Pro	Glu	Arg	Leu	Pro	Ile
545					550					555					560
Arg	Arg	Ser	Ser	Thr	Arg	Asp	Lys	Asn	Arg	Arg	Gly	Glu	Thr	Cys	Phe
				565					570					575	
Leu	Leu	Thr	Thr	Gly	Asp	Tyr	Thr	Cys	Ala	Ser	Asp	Gly	Gly	Ile	Arg
				580				585						590	
Lys	Val	Leu	Tyr	Arg	Ile	Tyr	His	Gly	Phe	Leu	Pro	Ala	Glu	Asn	Gly
	595						600					605			
Thr	Leu	Arg	Phe	Ser	His	Ser	Lys	Asp	Cys	Thr	Gly	Asn	Phe	Cys	Tyr
	610					615					620				

<210> 48  
 <211> 98  
 <212> PRT  
 <213> Homo sapiens

<400> 48															
Arg	Val	Arg	Leu	Asn	Val	Gly	Gly	Lys	Arg	Phe	Glu	Thr	Ser	Lys	Ser
1				5					10					15	
Thr	Leu	Thr	Arg	Phe	Pro	Asp	Thr	Arg	Leu	Gly	Arg	Leu	Leu	Glu	Cys
			20					25					30		
Arg	Asp	Cys	Asp	Phe	Tyr	Asp	Asp	Ala	Thr	Gly	Glu	Tyr	Phe	Phe	Asp
		35					40					45			
Arg	Ser	Pro	Lys	His	Phe	Glu	His	Ile	Leu	Asn	Phe	Tyr	Arg	Thr	Gly
	50					55					60				
Gly	Lys	Leu	His	Arg	Pro	Glu	Glu	Val	Cys	Leu	Glu	Ser	Phe	Leu	Glu
65					70					75					80
Glu	Leu	Glu	Phe	Tyr	Gly	Leu	Asp	Glu	Leu	Ala	Ile	Glu	Leu	Cys	Cys
				85					90					95	

Glu Asp

<210> 49  
 <211> 185

<212> PRT

<213> Homo sapiens

<400> 49

Leu Glu Ile Leu Asp Tyr Val Phe Thr Val Ile Phe Thr Leu Glu Met  
 1 5 10 15  
 Leu Leu Lys Phe Ile Ala Leu Gly Phe Lys Leu Lys Tyr Leu Arg Ser  
 20 25 30  
 Pro Trp Asn Ile Leu Asp Phe Leu Ile Val Leu Pro Ser Leu Ile Asp  
 35 40 45  
 Leu Ile Leu Phe Leu Ser Gly Gly Gly Ser Val Leu Arg Leu Leu Arg  
 50 55 60  
 Leu Leu Arg Leu Leu Arg Leu Leu Arg Arg Leu Glu Gly Leu Arg Thr  
 65 70 75 80  
 Leu Leu Gln Ser Leu Gly Arg Ser Leu Lys Ser Leu Leu Asn Leu Leu  
 85 90 95  
 Leu Leu Leu Leu Leu Leu Phe Ile Phe Ala Ile Ile Gly Val Gln  
 100 105 110  
 Leu Phe Gly Gly Glu Phe Asn Lys Cys Cys Asp Gly Val Asn Pro Ile  
 115 120 125  
 Asn Gly Asn Ser Asn Phe Asp Ser Phe Gly Glu Ala Phe Tyr Trp Leu  
 130 135 140  
 Phe Arg Thr Leu Thr Thr Glu Gly Trp Gly Asp Ile Met Pro Asp Thr  
 145 150 155 160  
 Leu Asp Ala Pro Val Leu Gly Lys Ile Phe Phe Val Ile Phe Ile Ile  
 165 170 175  
 Leu Gly Gly Leu Leu Leu Leu Asn Leu  
 180 185

<210> 50

<211> 95

<212> PRT

<213> Homo sapiens

<400> 50

Val Thr Leu Asn Val Gly Gly Lys Lys Phe His Ala His Lys Ala Val  
 1 5 10 15  
 Leu Ala Ala His Ser Pro Tyr Phe Lys Ala Leu Phe Ser Ser Asp Phe  
 20 25 30  
 Lys Glu Ser Asp Lys Ser Glu Ile Tyr Leu Phe Asp Val Ser Pro Glu  
 35 40 45  
 Asp Phe Arg Ala Leu Leu Asn Phe Leu Tyr Thr Gly Lys Leu Asp Ile



Gly Ser Gln Arg Ala Lys Arg Lys Val Thr Arg Met Ile Val Ile Val  
 225 230 235 240  
 Ala Val Leu Phe Cys Leu Cys Trp Met Pro His His Ala Leu Ile Leu  
 245 250 255  
 Cys Val Trp Phe Gly Arg Phe Pro Leu Thr Arg Ala Thr Tyr Ala Leu  
 260 265 270  
 Arg Ile Leu Ser His Leu Val Ser Tyr Ala Asn Ser Cys Val Asn Pro  
 275 280 285  
 Ile Val Tyr Ala Leu Val Ser Lys His Phe Arg Lys Gly Phe Arg Lys  
 290 295 300  
 Ile Cys Ala Gly Leu Leu Arg Arg Ala Pro Arg Arg Ala Ser Gly Arg  
 305 310 315 320  
 Val Cys Ile Leu Ala Pro Gly Asn His Ser Gly Gly Met Leu Glu Pro  
 325 330 335  
 Glu Ser Thr Asp Leu Thr Gln Val Ser Glu Ala Ala Gly Pro Leu Val  
 340 345 350  
 Pro Ala Pro Ala Leu Pro Asn Cys Thr Thr Leu Ser Arg Thr Leu Asp  
 355 360 365  
 Pro Ala Cys  
 370

<210> 52  
 <211> 387  
 <212> PRT  
 <213> Homo sapiens

<400> 52  
 Met Asn Val Ser Gly Cys Pro Gly Ala Gly Asn Ala Ser Gln Ala Gly  
 1 5 10 15  
 Gly Gly Gly Gly Trp His Pro Glu Ala Val Ile Val Pro Leu Leu Phe  
 20 25 30  
 Ala Leu Ile Phe Leu Val Gly Thr Val Gly Asn Thr Leu Val Leu Ala  
 35 40 45  
 Val Leu Leu Arg Gly Gly Gln Ala Val Ser Thr Thr Asn Leu Phe Ile  
 50 55 60  
 Leu Asn Leu Gly Val Ala Asp Leu Cys Phe Ile Leu Cys Cys Val Pro  
 65 70 75 80  
 Phe Gln Ala Thr Ile Tyr Thr Leu Asp Gly Trp Val Phe Gly Ser Leu  
 85 90 95  
 Leu Cys Lys Ala Val His Phe Leu Ile Phe Leu Thr Met His Ala Ser  
 100 105 110

Ser Phe Thr Leu Ala Ala Val Ser Leu Asp Arg Tyr Leu Ala Ile Arg  
115 120 125

Tyr Pro Leu His Ser Arg Glu Leu Arg Thr Pro Arg Asn Ala Leu Ala  
130 135 140

Ala Ile Gly Leu Ile Trp Gly Leu Ser Leu Leu Phe Ser Gly Pro Tyr  
145 150 155 160

Leu Ser Tyr Tyr Arg Gln Ser Gln Leu Ala Asn Leu Thr Val Cys His  
165 170 175

Pro Ala Trp Ser Ala Pro Arg Arg Arg Ala Met Asp Ile Cys Thr Phe  
180 185 190

Val Phe Ser Tyr Leu Leu Pro Val Leu Val Leu Gly Leu Thr Tyr Ala  
195 200 205

Arg Thr Leu Arg Tyr Leu Trp Arg Ala Val Asp Pro Val Ala Ala Gly  
210 215 220

Ser Gly Ala Arg Arg Ala Lys Arg Lys Val Thr Arg Met Ile Leu Ile  
225 230 235 240

Val Ala Ala Leu Phe Cys Leu Cys Trp Met Pro His His Ala Leu Ile  
245 250 255

Leu Cys Val Trp Phe Gly Gln Phe Pro Leu Thr Arg Ala Thr Tyr Ala  
260 265 270

Leu Arg Ile Leu Ser His Leu Val Ser Tyr Ala Asn Ser Cys Val Asn  
275 280 285

Pro Ile Val Tyr Ala Leu Val Ser Lys His Phe Arg Lys Gly Phe Arg  
290 295 300

Thr Ile Cys Ala Gly Leu Leu Gly Arg Ala Pro Gly Arg Ala Ser Gly  
305 310 315 320

Arg Val Cys Ala Ala Ala Arg Gly Thr His Ser Gly Ser Val Leu Glu  
325 330 335

Arg Glu Ser Ser Asp Leu Leu His Met Ser Glu Ala Ala Gly Ala Leu  
340 345 350

Arg Pro Cys Pro Gly Ala Ser Gln Pro Cys Ile Leu Glu Pro Cys Pro  
355 360 365

Gly Pro Ser Trp Gln Gly Pro Lys Ala Gly Asp Ser Ile Leu Thr Val  
370 375 380

Asp Val Ala  
385

<211> 372

<212> PRT

<213> Rattus norvegicus

<400> 53

Met Asn Gly Ser Gly Ser Gln Gly Ala Glu Asn Thr Ser Gln Glu Gly  
1 5 10 15  
Gly Ser Gly Gly Trp Gln Pro Glu Ala Val Leu Val Pro Leu Phe Phe  
20 25 30  
Ala Leu Ile Phe Leu Val Gly Thr Val Gly Asn Ala Leu Val Leu Ala  
35 40 45  
Val Leu Leu Arg Gly Gly Gln Ala Val Ser Thr Thr Asn Leu Phe Ile  
50 55 60  
Leu Asn Leu Gly Val Ala Asp Leu Cys Phe Ile Leu Cys Cys Val Pro  
65 70 75 80  
Phe Gln Ala Thr Ile Tyr Thr Leu Asp Asp Trp Val Phe Gly Ser Leu  
85 90 95  
Leu Cys Lys Ala Val His Phe Leu Ile Phe Leu Thr Met His Ala Ser  
100 105 110  
Ser Phe Thr Leu Ala Ala Val Ser Leu Asp Arg Tyr Leu Ala Ile Arg  
115 120 125  
Tyr Pro Leu His Ser Arg Glu Leu Arg Thr Pro Arg Asn Ala Leu Ala  
130 135 140  
Ala Ile Gly Leu Ile Trp Gly Leu Ala Leu Leu Phe Ser Gly Pro Tyr  
145 150 155 160  
Leu Ser Tyr Tyr Arg Gln Ser Gln Leu Ala Asn Leu Thr Val Cys His  
165 170 175  
Pro Ala Trp Ser Ala Pro Arg Arg Arg Ala Met Asp Leu Cys Thr Phe  
180 185 190  
Val Phe Ser Tyr Leu Leu Pro Val Leu Val Leu Ser Leu Thr Tyr Ala  
195 200 205  
Arg Thr Leu Arg Tyr Leu Trp Arg Thr Val Asp Pro Val Thr Ala Gly  
210 215 220  
Ser Gly Ser Gln Arg Ala Lys Arg Lys Val Thr Arg Met Ile Ile Ile  
225 230 235 240  
Val Ala Val Leu Phe Cys Leu Cys Trp Met Pro His His Ala Leu Ile  
245 250 255  
Leu Cys Val Trp Phe Gly Arg Phe Pro Leu Thr Arg Ala Thr Tyr Ala  
260 265 270  
Leu Arg Ile Leu Ser His Leu Val Ser Tyr Ala Asn Ser Cys Val Asn

275 280 285  
 Pro Ile Val Tyr Ala Leu Val Ser Lys His Phe Arg Lys Gly Phe Arg  
 290 295 300  
 Lys Ile Cys Ala Gly Leu Leu Arg Pro Ala Pro Arg Arg Ala Ser Gly  
 305 310 315 320  
 Arg Val Ser Ile Leu Ala Pro Gly Asn His Ser Gly Ser Met Leu Glu  
 325 330 335  
 Gln Glu Ser Thr Asp Leu Thr Gln Val Ser Glu Ala Ala Gly Pro Leu  
 340 345 350  
 Val Pro Pro Pro Ala Leu Pro Asn Cys Thr Ala Ser Ser Arg Thr Leu  
 355 360 365  
 Asp Pro Ala Cys  
 370

<210> 54  
 <211> 371  
 <212> PRT  
 <213> Mus musculus

<220>  
 <221> VARIANT  
 <222> (325)  
 <223> Wherein Xaa is any amino acid as defined in the  
 specification.

<220>  
 <221> VARIANT  
 <222> (360)  
 <223> Wherein Xaa is any amino acid as defined in the  
 specification

<400> 54  
 Met Asn Gly Ser Asp Ser Gln Gly Ala Glu Asp Ser Ser Gln Glu Gly  
 1 5 10 15  
 Gly Gly Gly Trp Gln Pro Glu Ala Val Leu Val Pro Leu Phe Phe Ala  
 20 25 30  
 Leu Ile Phe Leu Val Gly Ala Val Gly Asn Ala Leu Val Leu Ala Val  
 35 40 45  
 Leu Leu Arg Gly Gly Gln Ala Val Ser Thr Thr Asn Leu Phe Ile Leu  
 50 55 60  
 Asn Leu Gly Val Ala Asp Leu Cys Phe Ile Leu Cys Cys Val Pro Phe  
 65 70 75 80  
 Gln Ala Thr Ile Tyr Thr Leu Asp Asp Trp Val Phe Gly Ser Leu Leu  
 85 90 95



Cys Lys Ala Val His Phe Leu Ile Phe Leu Thr Met His Ala Ser Ser  
 100 105 110  
 Phe Thr Leu Ala Ala Val Ser Leu Asp Arg Tyr Leu Ala Ile Arg Tyr  
 115 120 125  
 Pro Leu His Ser Arg Glu Leu Arg Thr Pro Arg Asn Ala Leu Ala Ala  
 130 135 140  
 Ile Gly Leu Ile Trp Gly Leu Ala Leu Leu Phe Ser Gly Pro Tyr Leu  
 145 150 155 160  
 Ser Tyr Tyr Ser Gln Ser Gln Leu Ala Asn Leu Thr Val Cys His Pro  
 165 170 175  
 Ala Trp Ser Ala Pro Arg Arg Pro Trp Asn Ser Cys Thr Phe Cys Leu  
 180 185 190  
 Ser Tyr Leu Leu Pro Val Leu Val Leu Ser Leu Thr Tyr Ala Arg Thr  
 195 200 205  
 Leu His Tyr Leu Trp Arg Thr Val Asp Pro Val Val Ala Gly Ser Gly  
 210 215 220  
 Ser Gln Arg Ala Lys Arg Lys Val Thr Arg Met Ile Val Ile Val Ala  
 225 230 235 240  
 Val Leu Phe Cys Leu Cys Trp Met Pro His His Ala Leu Ile Leu Cys  
 245 250 255  
 Val Trp Phe Gly Arg Phe Pro Leu Thr Arg Ala Thr Tyr Ala Leu Arg  
 260 265 270  
 Ile Leu Ser His Leu Val Ser Tyr Ala Asn Ser Cys Val Asn Pro Ile  
 275 280 285  
 Val Tyr Ala Leu Val Ser Lys His Phe Arg Lys Gly Phe Arg Lys Ile  
 290 295 300  
 Cys Ala Gly Leu Leu Arg Arg Ala Pro Arg Arg Ala Ser Gly Arg Val  
 305 310 315 320  
 Cys Ile Leu Ala Xaa Gly Asn His Ser Gly Gly Met Leu Glu Pro Glu  
 325 330 335  
 Ser Thr Asp Leu Thr Gln Val Lys Arg Gly Ser Arg Ala Pro Arg Pro  
 340 345 350  
 Arg Thr Arg Thr Ser Gln Thr Xaa Thr Thr Leu Ser Arg Thr Leu Asp  
 355 360 365  
 Pro Ala Cys  
 370

<210> 55  
 <211> 370



Leu Val Tyr Ser Leu Ala Ser Arg His Phe Arg Ala Arg Phe Arg Arg  
290 295 300

Leu Trp Pro Cys Gly Arg Arg Arg His Arg His His His Arg Ala His  
305 310 315 320

Arg Ala Leu Arg Arg Val Gln Pro Ala Ser Ser Gly Pro Ala Gly Tyr  
325 330 335

Pro Gly Asp Ala Arg Pro Arg Gly Trp Ser Met Glu Pro Arg Gly Asp  
340 345 350

Ala Leu Arg Gly Gly Gly Glu Thr Arg Leu Thr Leu Ser Pro Arg Gly  
355 360 365

Pro Gln  
370

<210> 56  
<211> 205  
<212> PRT  
<213> Homo sapiens

<400> 56  
Trp Val Phe Gly Asp Ala Leu Cys Lys Leu Val Gly Ala Leu Phe Val  
1 5 10 15

Val Asn Gly Tyr Ala Ser Ile Leu Leu Thr Ala Ile Ser Ile Asp  
20 25 30

Arg Tyr Leu Ala Ile Val His Pro Leu Arg Tyr Arg Arg Ile Arg Thr  
35 40 45

Pro Arg Arg Ala Lys Val Leu Ile Leu Leu Val Trp Val Leu Ala Leu  
50 55 60

Leu Leu Ser Leu Pro Pro Leu Leu Phe Ser Trp Leu Arg Thr Val Glu  
65 70 75 80

Glu Gly Asn Thr Thr Val Cys Leu Ile Asp Phe Pro Glu Glu Ser Val  
85 90 95

Lys Arg Ser Tyr Val Leu Leu Ser Thr Leu Val Gly Phe Val Leu Pro  
100 105 110

Leu Leu Val Ile Leu Val Cys Tyr Thr Arg Ile Leu Arg Thr Leu Arg  
115 120 125

Lys Arg Ala Arg Ser Gln Arg Ser Leu Lys Arg Arg Ser Ser Ser Glu  
130 135 140

Arg Lys Ala Ala Lys Met Leu Leu Val Val Val Val Phe Val Leu  
145 150 155 160

Cys Trp Leu Pro Tyr His Ile Val Leu Leu Leu Asp Ser Leu Cys Leu

	165		170		175
Leu Ser Ile Trp Arg Val Leu Pro Thr Ala Leu Leu Ile Thr Leu Trp					
	180		185		190
Leu Ala Tyr Val Asn Ser Cys Leu Asn Pro Ile Ile Tyr					
	195		200		205
<210> 57					
<211> 337					
<212> PRT					
<213> Homo sapiens					
<400> 57					
Met Asn Glu Pro Leu Asp Tyr Leu Ala Asn Ala Ser Asp Phe Pro Asp					
1	5		10		15
Tyr Ala Ala Ala Phe Gly Asn Cys Thr Asp Glu Asn Ile Pro Leu Lys					
	20		25		30
Met His Tyr Leu Pro Val Ile Tyr Gly Ile Ile Phe Leu Val Gly Phe					
	35		40		45
Pro Gly Asn Ala Val Val Ile Ser Thr Tyr Ile Phe Lys Met Arg Pro					
	50		55		60
Trp Lys Ser Ser Thr Ile Ile Met Leu Asn Leu Ala Cys Thr Asp Leu					
	65		70		75
Leu Tyr Leu Thr Ser Leu Pro Phe Leu Ile His Tyr Tyr Ala Ser Gly					
		85	90		95
Glu Asn Trp Ile Phe Gly Asp Phe Met Cys Lys Phe Ile Arg Phe Ser					
	100		105		110
Phe His Phe Asn Leu Tyr Ser Ser Ile Leu Phe Leu Thr Cys Phe Ser					
	115		120		125
Ile Phe Arg Tyr Cys Val Ile Ile His Pro Met Ser Cys Phe Ser Ile					
	130		135		140
His Lys Thr Arg Cys Ala Val Val Ala Cys Ala Val Val Trp Ile Ile					
	145		150		155
Ser Leu Val Ala Val Ile Pro Met Thr Phe Leu Ile Thr Ser Thr Asn					
	165		170		175
Arg Thr Asn Arg Ser Ala Cys Leu Asp Leu Thr Ser Ser Asp Glu Leu					
	180		185		190
Asn Thr Ile Lys Trp Tyr Asn Leu Ile Leu Thr Ala Thr Thr Phe Cys					
	195		200		205
Leu Pro Leu Val Ile Val Thr Leu Cys Tyr Thr Thr Ile Ile His Thr					
	210		215		220

*(The following musical notation is transcribed from the handwritten manuscript as it appears in the document.)*

Phe His Ile Leu Arg Val Ile Arg Ile Glu Ser Arg Leu Leu Ser Ile  
260 265 270

Ser Cys Ser Ile Glu Asn Gln Ile His Glu Ala Tyr Ile Val Ser Arg  
275 280 285

Pro Leu Ala Ala Leu Asn Thr Phe Gly Asn Leu Leu Leu Tyr Val Val  
290 295 300

Val	Ser	Asp	Asn	Phe	Gln	Gln	Ala	Val	Cys	Ser	Thr	Val	Arg	Cys	Lys
305					310					315					320

Val Ser Gly Asn Leu Glu Gln Ala Lys Lys Ile Ser Tyr Ser Asn Asn  
325 330 335

Pro

```
<210> 58
<211> 373
<212> PRT
<213> Mus musculus
```

<400> 58  
Met Thr Glu Val Pro Trp Ser Val Val Pro Asn Gly Thr Asp Ala Ala  
1 5 10 15

Phe Leu Ala Gly Leu Gly Ser Leu Trp Gly Asn Ser Thr Val Ala Ser  
20 25 30

Thr Ala Ala Val Ser Ser Ser Phe Gln Cys Ala Leu Thr Lys Thr Gly  
35 40 45

Phe Gln Phe Tyr Tyr Leu Pro Ala Val Tyr Ile Leu Val Phe Ile Ile  
50 55 60

Gly Phe Leu Gly Asn Ser Val Ala Ile Trp Met Phe Val Phe His Met  
65 70 75 80

Lys Pro Trp Ser Gly Ile Ser Val Tyr Met Phe Asn Leu Ala Leu Ala  
85 90 95

Asp Phe Leu Tyr Val Leu Thr Leu Pro Ala Leu Ile Phe Tyr Tyr Phe  
100 105 110

Asn Lys Thr Asp Trp Ile Phe Gly Asp Ala Met Cys Lys Leu Gln Arg  
115 120 125

Phe Ile Phe His Val Asn Leu Tyr Gly Ser Ile Leu Phe Leu Thr Cys  
130 135 140



20					25					30					
Thr	Ala	Ala	Val	Ser	Ser	Ser	Phe	Lys	Cys	Ala	Leu	Thr	Lys	Thr	Gly
	35						40					45			
Phe	Gln	Phe	Tyr	Tyr	Leu	Pro	Ala	Val	Tyr	Ile	Leu	Val	Phe	Ile	Ile
	50					55					60				
Gly	Phe	Leu	Gly	Asn	Ser	Val	Ala	Ile	Trp	Met	Phe	Val	Phe	His	Met
	65					70					75				80
Lys	Pro	Trp	Ser	Gly	Ile	Ser	Val	Tyr	Met	Phe	Asn	Leu	Ala	Leu	Ala
				85					90					95	
Asp	Phe	Leu	Tyr	Val	Leu	Thr	Leu	Pro	Ala	Leu	Ile	Phe	Tyr	Tyr	Phe
			100					105					110		
Asn	Lys	Thr	Asp	Trp	Ile	Phe	Gly	Asp	Ala	Met	Cys	Lys	Leu	Gln	Arg
			115				120					125			
Phe	Ile	Phe	His	Val	Asn	Leu	Tyr	Gly	Ser	Ile	Leu	Phe	Leu	Thr	Cys
	130					135					140				
Ile	Ser	Ala	His	Arg	Tyr	Ser	Gly	Val	Val	Tyr	Pro	Leu	Lys	Ser	Leu
	145					150					155				160
Gly	Arg	Leu	Lys	Lys	Lys	Asn	Ala	Ile	Cys	Ile	Ser	Val	Leu	Val	Trp
				165					170					175	
Leu	Ile	Val	Val	Val	Ala	Ile	Ser	Pro	Ile	Leu	Phe	Tyr	Ser	Gly	Thr
			180					185					190		
Gly	Val	Arg	Lys	Asn	Lys	Thr	Ile	Thr	Cys	Tyr	Asp	Thr	Thr	Ser	Asp
		195					200					205			
Glu	Tyr	Leu	Arg	Ser	Tyr	Phe	Ile	Tyr	Ser	Met	Cys	Thr	Thr	Val	Ala
	210					215					220				
Met	Phe	Cys	Val	Pro	Leu	Val	Leu	Ile	Leu	Gly	Cys	Tyr	Gly	Leu	Ile
	225					230					235				240
Val	Arg	Ala	Leu	Ile	Tyr	Lys	Asp	Leu	Asp	Asn	Ser	Pro	Leu	Arg	Arg
				245					250					255	
Lys	Ser	Ile	Tyr	Leu	Val	Ile	Ile	Val	Leu	Thr	Val	Phe	Ala	Val	Ser
			260					265					270		
Tyr	Ile	Pro	Phe	His	Val	Met	Lys	Thr	Met	Asn	Leu	Arg	Ala	Arg	Leu
		275					280					285			
Asp	Phe	Gln	Thr	Pro	Ala	Met	Cys	Ala	Phe	Asn	Asp	Arg	Val	Tyr	Ala
	290					295					300				
Thr	Tyr	Gln	Val	Thr	Arg	Gly	Leu	Ala	Ser	Leu	Asn	Ser	Cys	Val	Asp
	305					310					315				320
Pro	Ile	Leu	Tyr	Phe	Leu	Ala	Gly	Asp	Thr	Phe	Arg	Arg	Arg	Leu	Ser





Glu Tyr Leu Arg Ser Tyr Phe Ile Tyr Ser Met Cys Thr Thr Val Ala  
 210 215 220

Met Phe Cys Ile Pro Leu Val Leu Ile Leu Gly Cys Tyr Gly Leu Ile  
 225 230 235 240

Val Arg Ala Leu Ile Tyr Lys Asp Leu Asp Asn Ser Pro Leu Arg Arg  
 245 250 255

Lys Ser Ile Tyr Leu Val Ile Ile Val Leu Thr Val Phe Ala Val Ser  
 260 265 270

Tyr Ile Pro Phe His Val Met Lys Thr Met Asn Leu Arg Ala Arg Leu  
 275 280 285

Asp Phe Gln Thr Pro Glu Met Cys Asp Phe Asn Asp Arg Val Tyr Ala  
 290 295 300

Thr Tyr Gln Val Thr Arg Gly Leu Ala Ser Leu Asn Ser Cys Val Asp  
 305 310 315 320

Pro Ile Leu Tyr Phe Leu Ala Gly Asp Thr Phe Arg Arg Arg Leu Ser  
 325 330 335

Arg Ala Thr Arg Lys Ala Ser Arg Arg Ser Glu Ala Asn Leu Gln Ser  
 340 345 350

Lys Ser Glu Glu Met Thr Leu Asn Ile Leu Ser Glu Phe Lys Gln Asn  
 355 360 365

Gly Asp Thr Ser Leu  
 370

<210> 61  
 <211> 373  
 <212> PRT  
 <213> Bos taurus

<400> 61  
 Met Thr Glu Val Leu Trp Pro Ala Val Pro Asn Gly Thr Asp Thr Ala  
 1 5 10 15

Phe Leu Ala Asp Pro Gly Ser Pro Trp Gly Asn Ser Thr Val Thr Ser  
 20 25 30

Thr Ala Ala Val Ala Ser Pro Phe Lys Cys Ala Leu Thr Lys Thr Gly  
 35 40 45

Phe Gln Phe Tyr Tyr Leu Pro Ala Val Tyr Ile Leu Val Phe Ile Ile  
 50 55 60

Gly Phe Leu Gly Asn Ser Val Ala Ile Trp Met Phe Val Phe His Met  
 65 70 75 80

Lys Pro Trp Ser Gly Ile Ser Val Tyr Met Phe Asn Leu Ala Leu Ala  
 85 90 95



1. The first group of people who are interested in the results of the study are the researchers themselves. They want to know if the study was successful in achieving its goals and if the data collected is reliable and valid. They also want to know if the study has contributed to the field of research and if it has provided any new insights or findings.

Gly Asn Leu Leu Val Ile Leu Val Ile Leu Arg Thr Lys Lys Leu Arg  
1 5 10 15

Phe Leu Leu Thr Leu Pro Pro Trp Ala Leu Tyr Tyr Leu Val Gly Gly  
35 40 45

Asp Trp Val Phe Gly Asp Ala Leu Cys Lys Leu Val Gly Ala Leu Phe  
50 55 60

Val Val Asn Gly Tyr Ala Ser Ile Leu Leu Leu Thr Ala Ile Ser Ile  
65 70 75 80

Asp Arg Tyr Leu Ala Ile Val His Pro Leu Arg Tyr Arg Arg Ile Arg  
85 90 95

Thr Pro Arg Arg Ala Lys Val Leu Ile Leu Leu Val Trp Val Leu Ala  
100 105 110

Leu Leu Leu Ser Leu Pro Pro Leu Leu Phe Ser Trp Leu Arg Thr Val  
115 120 125

Glu Glu Gly Asn Thr Thr Val Cys Leu Ile Asp Phe Pro Glu Glu Ser  
130 135 140

Val Lys Arg Ser Tyr Val Leu Leu Ser Thr Leu Val Gly Phe Val Leu  
145 150 155 160

Pro Leu Leu Val Ile Leu Val Cys Tyr Thr Arg Ile Leu Arg Thr Leu  
165 170 175

Arg Lys Arg Ala Arg Ser Gln Arg Ser Leu Lys Arg Arg Ser Ser Ser  
180 185 190

Glu Arg Lys Ala Ala Lys Met Leu Leu Val Val Val Val Val Phe Val  
195 200 205

Leu Cys Trp Leu Pro Tyr His Ile Val Leu Leu Leu Asp Ser Leu Cys  
210 215 220

Leu Leu Ser Ile Trp Arg Val Leu Pro Thr Ala Leu Leu Ile Thr Leu  
225 230 235 240

Trp Leu Ala Tyr Val Asn Ser Cys Leu Asn Pro Ile Ile Tyr  
245 250

79

<213> Homo sapiens

<400> 63

Met	Glu	Ser	Thr	Cys	Val	Ser	Ala	Ser	Leu	Pro	Arg	Ser	Tyr	Arg	Lys
1				5					10					15	
Thr	Asp	Thr	Val	Arg	Leu	Thr	Ser	Val	Val	Thr	Pro	Arg	Pro	Phe	Gly
			20					25					30		
Ser	Gln	Thr	Arg	Gly	Ile	Ser	Ser	Leu	Pro	Arg	Ser	Tyr	Thr	Met	Asp
		35					40					45			
Asp	Ala	Trp	Lys	Tyr	Asn	Gly	Asp	Ile	Glu	Asp	Ile	Lys	Arg	Thr	Pro
	50					55					60				
Asn	Asn	Val	Val	Ser	Thr	Pro	Ala	Pro	Ser	Pro	Asp	Ala	Ser	Gln	Leu
65					70					75					80
Ala	Ser	Ser	Leu	Ser	Ser	Gln	Lys	Glu	Val	Ala	Ala	Thr	Glu	Glu	Asp
				85					90					95	
Val	Thr	Arg	Leu	Pro	Ser	Pro	Thr	Ser	Pro	Phe	Ser	Ser	Leu	Ser	Gln
			100					105					110		
Asp	Gln	Ala	Ala	Thr	Ser	Lys	Ala	Thr	Leu	Ser	Ser	Thr	Ser	Gly	Leu
		115					120					125			
Asp	Leu	Met	Ser	Glu	Ser	Gly	Glu	Gly	Glu	Ile	Ser	Pro	Gln	Arg	Glu
	130					135					140				
Val	Ser	Arg	Ser	Gln	Asp	Gln	Phe	Ser	Asp	Met	Arg	Ile	Ser	Ile	Asn
145					150					155					160
Gln	Thr	Pro	Gly	Lys	Ser	Leu	Asp	Phe	Gly	Phe	Thr	Ile	Lys	Trp	Asp
				165					170					175	
Ile	Pro	Gly	Ile	Phe	Val	Ala	Ser	Val	Glu	Ala	Gly	Ser	Pro	Ala	Glu
			180					185					190		
Phe	Ser	Gln	Leu	Gln	Val	Asp	Asp	Glu	Ile	Ile	Ala	Ile	Asn	Asn	Thr
		195					200					205			
Lys	Phe	Ser	Tyr	Asn	Asp	Ser	Lys	Glu	Trp	Glu	Glu	Ala	Met	Ala	Lys
	210					215					220				
Ala	Gln	Glu	Thr	Gly	His	Leu	Val	Met	Asp	Val	Arg	Arg	Tyr	Gly	Lys
225					230					235					240
Ala	Gly	Ser	Pro	Glu	Thr	Lys	Trp	Ile	Asp	Ala	Thr	Ser	Gly	Ile	Tyr
				245					250					255	
Asn	Ser	Glu	Lys	Ser	Ser	Asn	Leu	Ser	Val	Thr	Thr	Asp	Phe	Ser	Glu
			260					265					270		
Ser	Leu	Gln	Ser	Ser	Asn	Ile	Glu	Ser	Lys	Glu	Ile	Asn	Gly	Ile	His
		275					280					285			

Asp Glu Ser Asn Ala Phe Glu Ser Lys Ala Ser Glu Ser Ile Ser Leu  
 290 295 300  
 Lys Asn Leu Lys Arg Arg Ser Gln Phe Phe Glu Gln Gly Ser Ser Asp  
 305 310 315 320  
 Ser Val Val Pro Asp Leu Pro Val Pro Thr Ile Ser Ala Pro Ser Arg  
 325 330 335  
 Trp Val Trp Asp Gln Glu Glu Glu Arg Lys Arg Gln Glu Arg Trp Gln  
 340 345 350  
 Lys Glu Gln Asp Arg Leu Leu Gln Glu Lys Tyr Gln Arg Glu Gln Glu  
 355 360 365  
 Lys Leu Arg Glu Glu Trp Gln Arg Ala Lys Gln Glu Ala Glu Arg Glu  
 370 375 380  
 Asn Ser Lys Tyr Leu Asp Glu Glu Leu Met Val Leu Ser Ser Asn Ser  
 385 390 395 400  
 Met Ser Leu Thr Thr Arg Glu Pro Ser Leu Ala Thr Trp Glu Ala Thr  
 405 410 415  
 Trp Ser Glu Gly Ser Lys Ser Ser Asp Arg Glu Gly Thr Arg Ala Gly  
 420 425 430  
 Glu Glu Glu Arg Arg Gln Pro Gln Glu Glu Val Val His Glu Asp Gln  
 435 440 445  
 Gly Lys Lys Pro Gln Asp Gln Leu Val Ile Glu Arg Glu Arg Lys Trp  
 450 455 460  
 Glu Gln Gln Leu Gln Glu Glu Gln Glu Gln Lys Arg Leu Gln Ala Glu  
 465 470 475 480  
 Ala Glu Glu Gln Lys Arg Pro Ala Glu Glu Gln Lys Arg Gln Ala Glu  
 485 490 495  
 Ile Glu Arg Glu Thr Ser Val Arg Ile Tyr Gln Tyr Arg Arg Pro Val  
 500 505 510  
 Asp Ser Tyr Asp Ile Pro Lys Thr Glu Glu Ala Ser Ser Gly Phe Leu  
 515 520 525  
 Pro Gly Asp Arg Asn Lys Ser Arg Ser Thr Thr Glu Leu Asp Asp Tyr  
 530 535 540  
 Ser Thr Asn Lys Asn Gly Asn Asn Lys Tyr Leu Asp Gln Ile Gly Asn  
 545 550 555 560  
 Thr Thr Ser Ser Gln Arg Arg Ser Lys Lys Glu Gln Val Pro Ser Gly  
 565 570 575  
 Ala Glu Leu Glu Arg Gln Gln Ile Leu Gln Glu Met Arg Lys Arg Thr  
 580 585 590

Pro Leu His Asn Asp Asn Ser Trp Ile Arg Gln Arg Ser Ala Ser Val  
595 600 605

Asn Lys Glu Pro Val Ser Leu Pro Gly Ile Met Arg Arg Gly Glu Ser  
610 615 620

Leu Asp Asn Leu Asp Ser Pro Arg Ser Asn Ser Trp Arg Gln Pro Pro  
625 630 635 640

Trp Leu Asn Gln Pro Thr Gly Phe Tyr Ala Ser Ser Ser Val Gln Asp  
645 650 655

Phe Ser Arg Pro Pro Pro Gln Leu Val Ser Thr Ser Asn Arg Ala Tyr  
660 665 670

Met Arg Asn Pro Ser Ser Ser Val Pro Pro Pro Ser Ala Gly Ser Val  
675 680 685

Lys Thr Ser Thr Thr Gly Val Ala Thr Thr Gln Ser Pro Thr Pro Arg  
690 695 700

Ser His Ser Pro Ser Ala Ser Gln Ser Gly Ser Gln Leu Arg Asn Arg  
705 710 715 720

Ser Val Ser Gly Lys Arg Ile Cys Ser Tyr Cys Asn Asn Ile Leu Gly  
725 730 735

Lys Gly Ala Ala Met Ile Ile Glu Ser Leu Gly Leu Cys Tyr His Leu  
740 745 750

His Cys Phe Lys Cys Val Ala Cys Glu Cys Asp Leu Gly Gly Ser Ser  
755 760 765

Ser Gly Ala Glu Val Arg Ile Arg Asn His Gln Leu Tyr Cys Asn Asp  
770 775 780

Cys Tyr Leu Arg Phe Lys Ser Gly Arg Pro Thr Ala Met  
785 790 795

<210> 64

<211> 797

<212> PRT

<213> Homo sapiens

<400> 64

Met Glu Ser Thr Arg Val Ser Ala Ser Leu Pro Arg Ser Tyr Arg Lys  
1 5 10 15

Thr Asp Thr Val Arg Leu Thr Ser Val Val Thr Pro Arg Pro Phe Gly  
20 25 30

Ser Gln Thr Arg Gly Ile Ser Ser Leu Pro Arg Ser Tyr Thr Met Asp  
35 40 45

Asp Ala Trp Lys Tyr Asn Gly Asp Val Glu Asp Ile Lys Arg Thr Pro  
50 55 60

Asn Asn Val Val Ser Thr Pro Ala Pro Ser Pro Asp Ala Ser Gln Leu  
 65 70 75 80  
 Ala Ser Ser Leu Ser Ser Gln Lys Glu Val Ala Ala Thr Glu Glu Asp  
 85 90 95  
 Val Thr Arg Leu Pro Ser Pro Thr Ser Pro Phe Ser Ser Leu Ser Gln  
 100 105 110  
 Asp Gln Ala Ala Thr Ser Lys Ala Thr Leu Ser Ser Thr Ser Gly Leu  
 115 120 125  
 Asp Leu Met Ser Glu Ser Gly Glu Gly Glu Ile Ser Pro Gln Arg Glu  
 130 135 140  
 Val Ser Arg Ser Gln Asp Gln Phe Ser Asp Met Arg Ile Ser Ile Asn  
 145 150 155 160  
 Gln Thr Pro Gly Lys Ser Leu Asp Phe Gly Phe Thr Ile Lys Trp Asp  
 165 170 175  
 Ile Pro Gly Ile Phe Val Ala Ser Val Glu Ala Gly Ser Pro Ala Glu  
 180 185 190  
 Phe Ser Gln Leu Gln Val Asp Asp Glu Ile Ile Ala Ile Asn Asn Thr  
 195 200 205  
 Lys Phe Ser Tyr Asn Asp Ser Lys Glu Trp Glu Glu Ala Met Ala Lys  
 210 215 220  
 Ala Gln Glu Thr Gly His Leu Val Met Asp Val Arg Arg Tyr Gly Lys  
 225 230 235 240  
 Ala Gly Ser Pro Glu Thr Lys Trp Ile Asp Ala Thr Ser Gly Ile Tyr  
 245 250 255  
 Asn Ser Glu Lys Ser Ser Asn Leu Ser Val Thr Thr Asp Phe Ser Glu  
 260 265 270  
 Ser Leu Gln Ser Ser Asn Ile Glu Ser Lys Glu Ile Asn Gly Ile His  
 275 280 285  
 Asp Glu Ser Asn Ala Phe Glu Ser Lys Ala Ser Glu Ser Ile Ser Leu  
 290 295 300  
 Lys Asn Leu Lys Arg Arg Ser Gln Phe Phe Glu Gln Gly Ser Ser Asp  
 305 310 315 320  
 Ser Val Val Pro Asp Leu Pro Val Pro Thr Ile Ser Ala Pro Ser Arg  
 325 330 335  
 Trp Val Trp Asp Gln Glu Glu Glu Arg Lys Arg Gln Glu Arg Trp Gln  
 340 345 350  
 Lys Glu Gln Asp Arg Leu Leu Gln Glu Lys Tyr Gln Arg Glu Gln Glu  
 355 360 365

[illegible]





130	135	140
Val Ser Arg Ser Gln Asp Gln Phe Ser Asp Met Arg Ile Ser Ile Asn		
145	150	155 160
Gln Thr Pro Gly Lys Ser Leu Asp Phe Gly Phe Thr Ile Lys Trp Asp		
	165	170 175
Ile Pro Gly Ile Phe Val Ala Ser Val Glu Ala Gly Ser Pro Ala Glu		
	180	185 190
Phe Ser Gln Leu Gln Val Asp Asp Glu Ile Ile Ala Ile Asn Asn Thr		
	195	200 205
Lys Phe Ser Tyr Asn Asp Ser Lys Glu Trp Glu Glu Ala Met Ala Lys		
	210	215 220
Ala Gln Glu Thr Gly His Leu Val Met Asp Val Arg Arg Tyr Gly Lys		
	225	230 235 240
Ala Gly Ser Pro Glu Thr Lys Trp Ile Asp Ala Thr Ser Gly Ile Tyr		
	245	250 255
Asn Ser Glu Lys Ser Ser Asn Leu Ser Val Thr Thr Asp Phe Ser Glu		
	260	265 270
Ser Leu Gln Ser Ser Asn Ile Glu Ser Lys Glu Ile Asn Gly Ile His		
	275	280 285
Asp Glu Ser Asn Ala Phe Glu Ser Lys Ala Ser Glu Ser Ile Ser Leu		
	290	295 300
Lys Asn Leu Lys Arg Arg Ser Gln Phe Phe Glu Gln Gly Ser Ser Asp		
	305	310 315 320
Ser Val Val Pro Asp Leu Pro Val Pro Thr Ile Ser Ala Pro Ser Arg		
	325	330 335
Trp Val Trp Asp Gln Glu Glu Glu Arg Lys Arg Gln Glu Arg Trp Gln		
	340	345 350
Lys Glu Gln Asp Arg Leu Leu Gln Glu Lys Tyr Gln Arg Glu Gln Glu		
	355	360 365
Lys Leu Arg Glu Glu Trp Gln Arg Ala Lys Gln Glu Ala Glu Arg Glu		
	370	375 380
Asn Ser Lys Tyr Leu Asp Glu Glu Leu Met Val Leu Ser Ser Asn Ser		
	385	390 395 400
Met Ser Leu Thr Thr Arg Glu Pro Ser Leu Ala Thr Trp Glu Ala Thr		
	405	410 415
Trp Ser Glu Gly Ser Lys Ser Ser Asp Arg Glu Gly Thr Arg Ala Gly		
	420	425 430
Glu Glu Glu Arg Arg Gln Pro Gln Glu Glu Val Val His Glu Asp Gln		

435 440 445

Gly Lys Lys Pro Gln Asp Gln Leu Val Ile Glu Arg Glu Arg Lys Trp  
450 455 460

Glu Gln Gln Leu Gln Glu Glu Gln Glu Gln Lys Arg Leu Gln Ala Glu  
465 470 475 480

Ala Glu Glu Gln Lys Arg Pro Ala Glu Glu Gln Lys Arg Gln Ala Glu  
485 490 495

Ile Glu Arg Glu Thr Ser Val Arg Ile Tyr Gln Tyr Arg Arg Pro Val  
500 505 510

Asp Ser Tyr Asp Ile Pro Lys Thr Glu Glu Ala Ser Ser Gly Phe Leu  
515 520 525

Pro Gly Asp Arg Asn Lys Ser Arg Ser Thr Thr Glu Leu Asp Asp Tyr  
530 535 540

Ser Thr Asn Lys Asn Gly Asn Asn Lys Tyr Leu Asp Gln Ile Gly Asn  
545 550 555 560

Thr Thr Ser Ser Gln Arg Arg Ser Lys Lys Glu Gln Val Pro Ser Gly  
565 570 575

Ala Glu Leu Glu Arg Gln Gln Ile Leu Gln Glu Met Arg Lys Arg Thr  
580 585 590

Pro Leu His Asn Asp Asn Ser Trp Ile Arg Gln Arg Ser Ala Ser Val  
595 600 605

Asn Lys Glu Pro Val Ser Leu Pro Gly Ile Met Arg Arg Gly Glu Ser  
610 615 620

Leu Asp Asn Leu Asp Ser Pro Arg Ser Asn Ser Trp Arg Gln Pro Pro  
625 630 635 640

Trp Leu Asn Gln Pro Thr Gly Phe Tyr Ala Ser Ser Ser Val Gln Asp  
645 650 655

Phe Ser Arg Pro Pro Pro Gln Leu Val Ser Thr Ser Asn Arg Ala Tyr  
660 665 670

Met Arg Asn Pro Ser Ser Ser Val Pro Pro Pro Ser Ala Gly Ser Val  
675 680 685

Lys Thr Ser Thr Thr Gly Val Ala Thr Thr Gln Ser Pro Thr Pro Arg  
690 695 700

Ser His Ser Pro Ser Ala Ser Gln Ser Gly Ser Gln Leu Arg Asn Ser  
705 710 715 720

Val Leu Pro Val Ser Val Thr Ser Glu Ala Leu Pro Gln Glu Leu Lys  
725 730 735

Ser Gly Ser Glu Thr Thr Asn Cys Thr Ala Thr Thr Ala Ile Ser Asp













Phe Ser Arg Pro Pro Pro Gln Leu Val Ser Thr Ser Asn Arg Ala Tyr  
660 665 670

Met Arg Asn Pro Ser Ser Ser Val Pro Pro Pro Ser Ala Gly Ser Val  
675 680 685

Lys Thr Ser Thr Thr Gly Val Ala Thr Thr Gln Ser Pro Thr Pro Arg  
690 695 700

Ser His Ser Pro Ser Ala Ser Gln Ser Gly Ser Gln Leu Arg Asn Ser  
705 710 715 720

Val Leu Pro Val Ser Val Thr Ser Glu Ala Leu Pro Gln Glu Leu Lys  
725 730 735

Ser Gly Ser Glu Thr Thr Asn Cys Thr Ala Thr Thr Ala Ile Ser Asp  
740 745 750

Ser Asn Leu Asp Gly Gln Pro Pro Cys Asp Val Ser Leu His Thr Lys  
755 760 765

Ala Leu Leu Gln Ile Glu Glu Glu Val Val Ala Ala His Val Asp Leu  
770 775 780

<210> 68  
<211> 71  
<212> PRT  
<213> Homo sapiens

<400> 68  
Leu Gly Phe Ser Leu Val Gly Gly Lys Asp Ser Gly Asp Gly Gly Val  
1 5 10 15

Val Val Ser Ser Val Val Pro Gly Ser Pro Ala Ala Lys Ala Gly Leu  
20 25 30

Lys Pro Gly Asp Val Ile Leu Glu Val Asn Gly Thr Ser Val Glu Gly  
35 40 45

Leu Thr His Leu Glu Ala Val Asp Leu Leu Lys Glu Ala Gly Gly Lys  
50 55 60

Val Thr Leu Thr Val Leu Arg  
65 70

<210> 69  
<211> 561  
<212> PRT  
<213> Mus musculus

<400> 69

Met Ala Val Leu Leu Ala Ala Val Leu Ala Ser Ser Leu Tyr Leu Gln  
1 5 10 15

Val Ala Ala Asp Phe Asp Gly Arg Trp Pro Arg Gln Ile Val Ser Ser  
20 25 30

Ile Gly Leu Cys Arg Tyr Gly Gly Arg Ile Asp Cys Cys Trp Gly Trp  
35 40 45

Ala Arg Gln Ser Trp Gly Gln Cys Gln Pro Val Cys Gln Pro Gln Cys  
50 55 60

Lys His Gly Glu Cys Val Gly Pro Asn Lys Cys Lys Cys His Pro Gly  
65 70 75 80

Phe Ala Gly Lys Thr Cys Asn Gln Asp Leu Asn Glu Cys Gly Leu Lys  
85 90 95

Pro Arg Pro Cys Lys His Arg Cys Met Asn Thr Phe Gly Ser Tyr Lys  
100 105 110

Cys Tyr Cys Leu Asn Gly Tyr Met Leu Leu Pro Asp Gly Ser Cys Ser  
115 120 125

Ser Ala Leu Ser Cys Ser Met Ala Asn Cys Gln Tyr Gly Cys Asp Val  
130 135 140

Val Lys Gly Gln Val Arg Cys Gln Cys Pro Ser Pro Gly Leu Gln Leu  
145 150 155 160

Ala Pro Asp Gly Arg Thr Cys Val Asp Ile Asp Glu Cys Ala Thr Gly  
165 170 175

Arg Val Ser Cys Pro Arg Phe Arg Gln Cys Val Asn Thr Phe Gly Ser  
180 185 190

Tyr Ile Cys Lys Cys His Thr Gly Phe Asp Leu Met Tyr Ile Gly Gly  
195 200 205

Lys Tyr Gln Cys His Asp Ile Asp Glu Cys Ser Leu Gly Gln His Gln  
210 215 220

Cys Ser Ser Tyr Ala Arg Cys Tyr Asn Ile His Gly Ser Tyr Lys Cys  
225 230 235 240

Gln Cys Arg Asp Gly Tyr Glu Gly Asp Gly Leu Asn Cys Val Tyr Ile  
245 250 255

Pro Lys Val Met Ile Glu Pro Ser Gly Pro Ile His Met Pro Glu Arg  
260 265 270

Asn Gly Thr Ile Ser Lys Gly Asp Gly Gly His Ala Asn Arg Ile Pro  
275 280 285

Asp Ala Gly Ser Thr Arg Trp Pro Leu Lys Thr Pro Tyr Ile Pro Pro  
290 295 300





Pro Asp Ala Gly Ser Thr Arg Trp Pro Leu Lys Thr Pro Tyr Ile Pro  
305 310 315 320

Pro Val Ile Thr Asn Arg Pro Thr Ser Lys Pro Thr Thr Arg Pro Thr  
325 330 335

Pro Asn Pro Thr Pro Gln Pro Thr Pro Pro Pro Pro Pro Pro Leu Pro  
340 345 350

Thr Glu Pro Arg Thr Thr Pro Leu Pro Pro Thr Pro Glu Arg Pro Ser  
355 360 365

Thr Arg Pro Thr Thr Ile Ala Pro Ala Thr Ser Thr Thr Thr Arg Val  
370 375 380

Ile Thr Val Asp Asn Arg Ile Gln Thr Asp Pro Gln Lys Pro Arg Gly  
385 390 395 400

Asp Val Phe Ile Pro Arg Gln Pro Thr Asn Asp Leu Phe Glu Ile Phe  
405 410 415

Glu Ile Glu Arg Gly Val Ser Ala Asp Glu Glu Val Lys Asp Asp Pro  
420 425 430

Gly Ile Leu Ile His Ser Cys Asn Phe Asp His Gly Leu Cys Gly Trp  
435 440 445

Ile Arg Glu Lys Asp Ser Asp Leu His Trp Glu Thr Ala Arg Asp Pro  
450 455 460

Ala Gly Gly Gln Tyr Leu Thr Val Ser Ala Ala Lys Ala Pro Gly Gly  
465 470 475 480

Lys Ala Ala Arg Leu Val Leu Arg Leu Gly His Leu Met His Ser Gly  
485 490 495

Asp Leu Cys Leu Ser Phe Arg His Lys Val Thr Gly Leu His Ser Gly  
500 505 510

Thr Leu Gln Val Phe Val Arg Lys His Gly Thr His Gly Ala Ala Leu  
515 520 525

Trp Gly Arg Asn Gly Gly His Gly Trp Arg Gln Thr Gln Ile Thr Leu  
530 535 540

Arg Gly Ala Asp Val Lys Ser Val Ile Phe Lys Gly Glu Lys Arg Arg  
545 550 555 560

Gly His Thr Gly Glu Ile Gly Leu Asp Asp Val Ser Leu Lys Arg Gly  
565 570 575

Arg Cys









Tyr Ile Cys Lys Cys His Thr Gly Phe Asp Leu Met Tyr Ile Gly Gly  
 245 250 255  
 Lys Tyr Gln Cys His Asp Ile Asp Glu Cys Ser Leu Gly Gln His Gln  
 260 265 270  
 Cys Ser Ser Tyr Ala Arg Cys Tyr Asn Ile His Gly Ser Tyr Lys Cys  
 275 280 285  
 Gln Cys Arg Asp Gly Tyr Glu Gly Asp Gly Leu Asn Cys Val Tyr Ile  
 290 295 300  
 Pro Lys Val Met Ile Glu Pro Ser Gly Pro Ile His Met Pro Glu Arg  
 305 310 315 320  
 Asn Gly Thr Ile Ser Lys Gly Asp Gly Gly His Ala Asn Arg Ile Pro  
 325 330 335  
 Asp Ala Gly Ser Thr Arg Trp Pro Leu Lys Thr Pro Tyr Ile Pro Pro  
 340 345 350  
 Val Ile Thr Asn Arg Pro Thr Ser Lys Pro Thr Thr Arg Pro Thr Pro  
 355 360 365  
 Asn Pro Thr Pro Gln Pro Thr Pro Pro Pro Pro Pro Pro Leu Pro Thr  
 370 375 380  
 Glu Pro Arg Thr Thr Pro Leu Pro Pro Thr Pro Glu Arg Pro Ser Thr  
 385 390 395 400  
 Arg Pro Thr Thr Ile Ala Pro Ala Thr Ser Thr Thr Thr Arg Val Ile  
 405 410 415  
 Thr Val Asp Asn Arg Ile Gln Thr Asp Pro Gln Lys Pro Arg Gly Asp  
 420 425 430  
 Val Phe Ile Pro Arg Gln Pro Thr Asn Asp Leu Phe Glu Ile Phe Glu  
 435 440 445  
 Ile Glu Arg Gly Val Ser Ala Asp Glu Glu Val Lys Asp Asp Pro Gly  
 450 455 460  
 Ile Leu Ile His Ser Cys Asn Phe Asp His Gly Leu Cys Gly Trp Ile  
 465 470 475 480  
 Arg Glu Lys Asp Ser Asp Leu His Trp Glu Thr Ala Arg Asp Pro Ala  
 485 490 495  
 Gly Gly Gln Tyr Leu Thr Val Ser Ala Ala Lys Ala Pro Gly Gly Lys  
 500 505 510  
 Ala Ala Arg Leu Val Leu Arg Leu Gly His Leu Met His Ser Gly Asp  
 515 520 525  
 Leu Cys Leu Ser Phe Arg His Lys Val Thr Gly Leu His Ser Gly Thr  
 530 535 540



Phe Gly Ser Tyr Tyr Cys Lys Cys His Ile Gly Phe Glu Leu Lys Tyr  
195 200 205  
Ile Gly Arg Arg Tyr Asp Cys Val Asp Ile Asn Glu Cys Ala Leu Asn  
210 215 220  
Thr His Pro Cys Ser Pro His Ala Asn Cys Leu Asn Thr Arg Gly Ser  
225 230 235 240  
Phe Lys Cys Lys Cys Lys Gln Gly Tyr Arg Gly Asn Gly Leu Gln Cys  
245 250 255  
Ser Val Ile Pro Glu His Ser Val Lys Glu Ile Leu Thr Ala Pro Gly  
260 265 270  
Thr Ile Lys Asp Arg Ile Lys Lys Leu Leu Ala His Lys Arg Thr Met  
275 280 285  
Lys Lys Lys Val Lys Leu Lys Met Val Thr Pro Arg Pro Ala Ser Thr  
290 295 300  
Arg Val Pro Lys Val Asn Leu Pro Tyr Ser Ser Glu Glu Gly Val Ser  
305 310 315 320  
Arg Gly Arg Asn Tyr Asp Gly Glu Gln Lys Lys Lys Glu Glu Gly Lys  
325 330 335  
Arg Glu Arg Leu Glu Glu Glu Lys Gly Glu Lys Thr Leu Arg Asn Glu  
340 345 350  
Val Glu Gln Glu Arg Thr Leu Arg Gly Asp Val Phe Ser Pro Lys Val  
355 360 365  
Asn Glu Ala Glu Asp Leu Asp Leu Val Tyr Val Gln Arg Lys Glu Leu  
370 375 380  
Asn Ser Lys Leu Lys His Lys Asp Leu Asn Ile Ser Val Asp Cys Ser  
385 390 395 400  
Phe Asp Leu Gly Val Cys Asp Trp Lys Gln Asp Arg Glu Asp Asp Phe  
405 410 415  
Asp Trp His Pro Ala Asp Arg Asp Asn Asp Val Gly Tyr Tyr Met Ala  
420 425 430  
Val Pro Ala Leu Ala Gly His Lys Lys Asn Ile Gly Arg Leu Lys Leu  
435 440 445  
Leu Leu Pro Asn Leu Thr Pro Gln Ser Asn Phe Cys Leu Leu Phe Asp  
450 455 460  
Tyr Arg Leu Ala Gly Asp Lys Val Gly Lys Leu Arg Val Phe Val Lys  
465 470 475 480  
Asn Ser Asn Asn Ala Leu Ala Trp Glu Glu Thr Lys Asn Glu Asp Gly  
485 490 495

Arg Trp Arg Thr Gly Lys Ile Gln Leu Tyr Gln Gly Ile Asp Thr Thr  
500 505 510

Lys Ser Val Ile Phe Glu Ala Glu Arg Gly Lys Gly Lys Thr Gly Glu  
515 520 525

Ile Ala Val Asp Gly Val Leu Leu Val Ser Gly Leu Cys Pro Asp Asp  
530 535 540

Phe Leu Ser Val Glu Gly  
545 550

<210> 74  
<211> 158  
<212> PRT  
<213> Homo sapiens

<400> 74  
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Ser Asn Asp Asp Gly Pro Trp Glu Arg Val Ser Ser Ala Thr Arg Asn  
20 25 30

Asp Gly Pro Asn Arg Asp His Thr Thr Gly Asn Gly His Tyr Met Phe  
35 40 45

Phe Glu Thr Ser Ser Gly Lys Pro Gly Gln Thr Ala Arg Leu Leu Ser  
50 55 60

Pro Pro Leu Tyr Glu Asn Arg Ser Thr His Cys Leu Thr Phe Trp Tyr  
65 70 75 80

Tyr Met Tyr Gly Ser Gly Val Gly Thr Leu Asn Val Tyr Val Arg Val  
85 90 95

Asn Asn Gly Pro Gln Asp Thr Leu Leu Trp Ser Arg Ser Gly Thr Gln  
100 105 110

Gly Gly Gln Trp Leu Gln Ala Glu Val Ala Leu Ser Thr Ser Pro Gln  
115 120 125

Pro Phe Gln Val Val Phe Glu Gly Thr Arg Gly Gly Gly Pro Ser Gly  
130 135 140

Tyr Ile Ala Leu Asp Asp Ile Leu Leu Ser Asn Gly Pro Cys  
145 150 155

<210> 75  
<211> 159  
<212> PRT  
<213> Homo sapiens

<400> 75











Lys Arg Glu Thr Lys Met Arg Gly Met Met Ala Lys Gln Glu Gly Met  
 180 185 190  
 Glu Met Lys Leu Gln Val Thr Gln Arg Ser Leu Glu Glu Ser Gln Gly  
 195 200 205  
 Lys Ile Ala Gln Leu Glu Gly Lys Leu Val Ser Ile Glu Lys Glu Lys  
 210 215 220  
 Ile Asp Glu Lys Ser Glu Thr Glu Lys Leu Leu Glu Tyr Ile Glu Glu  
 225 230 235 240  
 Ile Ser Cys Ala Ser Asp Gln Val Glu Lys Tyr Lys Leu Asp Ile Ala  
 245 250 255  
 Gln Leu Glu Glu Asn Leu Lys Glu Lys Asn Asp Glu Ile Leu Ser Leu  
 260 265 270  
 Lys Gln Ser Leu Glu Asp Asn Ile Val Ile Leu Ser Lys Gln Val Glu  
 275 280 285  
 Asp Leu Asn Val Lys Cys Gln Leu Leu Glu Thr Glu Lys Glu Asp His  
 290 295 300  
 Val Asn Arg Asn Arg Glu His Asn Glu Asn Leu Asn Ala Glu Met Gln  
 305 310 315 320  
 Asn Leu Glu Gln Lys Phe Ile Leu Glu Gln Arg Glu His Glu Lys Leu  
 325 330 335  
 Gln Gln Lys Glu Leu Gln Ile Asp Ser Leu Leu Gln Gln Glu Lys Glu  
 340 345 350  
 Leu Ser Ser Ser Leu His Gln Lys Leu Cys Ser Phe Gln Glu Glu Met  
 355 360 365  
 Val Lys Glu Lys Asn Leu Phe Glu Glu Glu Leu Lys Gln Thr Leu Asp  
 370 375 380  
 Glu Leu Asp Lys Leu Gln Gln Lys Glu Glu Gln Ala Glu Arg Leu Val  
 385 390 395 400  
 Lys Gln Leu Glu Glu Glu Ala Lys Ser Arg Ala Glu Glu Leu Lys Leu  
 405 410 415  
 Leu Glu Glu Lys Leu Lys Gly Lys Glu Ala Glu Leu Glu Lys Ser Ser  
 420 425 430  
 Ala Ala His Thr Gln Ala Thr Leu Leu Leu Gln Glu Lys Tyr Asp Ser  
 435 440 445  
 Met Val Gln Ser Leu Glu Asp Val Thr Ala Gln Phe Glu Ser Tyr Lys  
 450 455 460  
 Ala Leu Thr Ala Ser Glu Ile Glu Asp Leu Lys Leu Glu Asn Ser Ser  
 465 470 475 480



Cys Ala Pro Ser Pro Gly Ala Tyr Asp Val Lys Thr Leu Glu Val Leu  
20 25 30

Lys Gly Pro Val Ser Phe Gln Lys Ser Gln Arg Phe Lys Gln Gln Lys  
35 40 45

Glu Ser Lys Gln Asn Leu Asn Val Asp Lys Asp Thr Thr Leu Pro Ala  
50 55 60

Ser Ala Arg Lys Val Lys Ser Ser Glu Ser Lys Lys Glu Ser Gln Lys  
65 70 75 80

Asn Asp Lys Asp Leu Lys Ile Leu Glu Lys Glu Ile Arg Val Leu Leu  
85 90 95

Gln Glu Arg Gly Ala Gln Asp Arg Arg Ile Gln Asp Leu Glu Thr Glu  
100 105 110

Leu Glu Lys Met Glu Ala Arg Leu Asn Ala Ala Leu Arg Glu Lys Thr  
115 120 125

Ser Leu Ser Ala Asn Asn Ala Thr Leu Glu Lys Gln Leu Ile Glu Leu  
130 135 140

Thr Arg Thr Asn Glu Leu Leu Lys Ser Lys Phe Ser Glu Asn Gly Asn  
145 150 155 160

Gln Lys Asn Leu Arg Ile Leu Ser Leu Glu Leu Met Lys Leu Arg Asn  
165 170 175

Lys Arg Glu Thr Lys Met Arg Gly Met Met Ala Lys Gln Glu Gly Met  
180 185 190

Glu Met Lys Leu Gln Val Thr Gln Arg Ser Leu Glu Glu Ser Gln Gly  
195 200 205

Lys Ile Ala Gln Leu Glu Gly Lys Leu Val Ser Ile Glu Lys Glu Lys  
210 215 220

Ile Asp Glu Lys Ser Glu Thr Glu Lys Leu Leu Glu Tyr Ile Glu Glu  
225 230 235 240

Ile Ser Cys Ala Ser Asp Gln Val Glu Lys Tyr Lys Leu Asp Ile Ala  
245 250 255

Gln Leu Glu Glu Asn Leu Lys Glu Lys Asn Asp Glu Ile Leu Ser Leu  
260 265 270

Lys Gln Ser Leu Glu Glu Asn Ile Val Ile Leu Ser Lys Gln Val Glu  
275 280 285

Asp Leu Asn Val Lys Cys Gln Leu Leu Glu Lys Glu Lys Glu Asp His  
290 295 300

Val Asn Arg Asn Arg Glu His Asn Glu Asn Leu Asn Ala Glu Met Gln  
305 310 315 320

Asn Leu Lys Gln Lys Phe Ile Leu Glu Gln Gln Glu His Glu Lys Leu  
325 330 335

Gln Gln Lys Glu Leu Gln Ile Asp Ser Leu Leu Gln Gln Glu Lys Glu  
340 345 350

Leu Ser Ser Ser Leu His Gln Lys Leu Cys Ser Phe Gln Glu Glu Met  
355 360 365

Val Lys Glu Lys Asn Leu Phe Glu Glu Glu Leu Lys Gln Thr Leu Asp  
370 375 380

Glu Leu Asp Lys Leu Gln Gln Lys Glu Glu Gln Ala Glu Arg Leu Val  
385 390 395 400

Lys Gln Leu Glu Glu Glu Ala Lys Ser Arg Ala Glu Glu Leu Lys Leu  
405 410 415

Leu Glu Glu Lys Leu Lys Gly Lys Glu Ala Glu Leu Glu Lys Ser Ser  
420 425 430

Ala Ala His Thr Gln Ala Thr Leu Leu Leu Gln Glu Lys Tyr Asp Ser  
435 440 445

Met Val Gln Ser Leu Glu Asp Val Thr Ala Gln Phe Glu Ser Tyr Lys  
450 455 460

Ala Leu Thr Ala Ser Glu Ile Glu Asp Leu Lys Leu Glu Asn Ser Ser  
465 470 475 480

Leu Gln Glu Lys Ala Ala Lys Ala Gly Lys Asn Ala Glu Asp Val Gln  
485 490 495

His Gln Ile Leu Ala Thr Glu Ser Ser Asn Gln Glu Tyr Val Arg Met  
500 505 510

Leu Leu Asp Leu Gln Thr Lys Ser Ala Leu Lys Glu Thr Glu Ile Lys  
515 520 525

Glu Ile Thr Val Ser Phe Leu Gln Lys Ile Thr Asp Leu Gln Asn Gln  
530 535 540

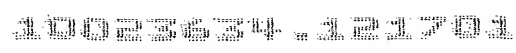
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545 550 555 560

Gly Arg Lys Ala Glu Lys Glu Asn Thr Thr Ala Glu Leu Thr Glu Glu  
565 570 575

Ile Asn Lys Trp Arg Leu Leu Tyr Glu Glu Leu Tyr Asn Lys Thr Lys  
580 585 590

Pro Phe Gln Leu Gln Leu Asp Ala Phe Glu Val Glu Lys Gln Ala Leu  
595 600 605

Leu Asn Glu His Gly Ala Ala Gln Glu Gln Leu Asn Lys Ile Arg Asp  
610 615 620



Ser Tyr Ala Lys Leu Leu Gly His Gln Asn Leu Lys Gln Lys Ile Lys  
625 630 635 640

His Val Val Lys Leu Lys Asp Glu Asn Ser Gln Leu Lys Ser Glu Val  
645 650 655

Ser Lys Leu Arg Cys Gln Leu Ala Lys Lys Lys Gln Ser Glu Thr Lys  
660 665 670

Leu Gln Glu Glu Leu Asn Lys Val Leu Gly Ile Lys His Phe Asp Pro  
675 680 685

Ser Lys Ala Phe His His Glu Ser Lys Glu Asn Phe Ala Leu Lys Thr  
690 695 700

Pro Leu Lys Glu Gly Asn Thr Asn Cys Tyr Arg Ala Pro Met Glu Cys  
705 710 715 720

Gln Glu Ser Trp Lys  
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<210> 80

<211> 724

<212> PRT

<213> Homo sapiens

<400> 80

Met Ser Phe Pro Lys Ala Pro Leu Lys Arg Phe Asn Asp Pro Ser Gly  
1 5 10 15

Cys Ala Pro Ser Pro Gly Ala Tyr Asp Val Lys Thr Leu Glu Val Leu  
20 25 30

Lys Gly Pro Val Ser Phe Gln Lys Ser Gln Arg Phe Lys Gln Gln Lys  
35 40 45

Glu Ser Lys Gln Asn Leu Asn Val Asp Lys Asp Thr Thr Leu Pro Ala  
50 55 60

Ser Ala Arg Lys Val Lys Ser Ser Glu Ser Lys Glu Ser Gln Lys Asn  
65 70 75 80

Asp Lys Asp Leu Lys Ile Leu Glu Lys Glu Ile Arg Val Leu Leu Gln  
85 90 95

Glu Arg Gly Ala Gln Asp Ser Arg Ile Gln Asp Leu Glu Thr Glu Leu  
100 105 110

Glu Lys Met Glu Ala Arg Leu Asn Ala Ala Leu Arg Glu Lys Thr Ser  
115 120 125

Leu Ser Ala Asn Asn Ala Thr Leu Glu Lys Gln Leu Ile Glu Leu Thr  
130 135 140

Arg Thr Asn Glu Leu Leu Lys Ser Lys Phe Ser Glu Asn Gly Asn Gln

145                      150                      155                      160

Lys Asn Leu Arg Ile Leu Ser Leu Glu Leu Met Lys Leu Arg Asn Lys  
                                  165                      170                      175

Arg Glu Thr Lys Met Arg Gly Met Met Ala Lys Gln Glu Gly Met Glu  
                                  180                      185                      190

Met Lys Leu Gln Val Thr Gln Arg Ser Leu Glu Glu Ser Gln Gly Lys  
                                  195                      200                      205

Ile Ala Gln Leu Glu Gly Lys Leu Val Ser Ile Glu Lys Glu Lys Ile  
                                  210                      215                      220

Asp Glu Lys Ser Glu Thr Glu Lys Leu Leu Glu Tyr Ile Glu Glu Ile  
                                  225                      230                      235                      240

Ser Cys Ala Ser Asp Gln Val Glu Lys Tyr Lys Leu Asp Ile Ala Gln  
                                  245                      250                      255

Leu Glu Glu Asn Leu Lys Glu Lys Asn Asp Glu Ile Leu Ser Leu Lys  
                                  260                      265                      270

Gln Ser Leu Glu Glu Asn Ile Val Ile Leu Ser Lys Gln Val Glu Asp  
                                  275                      280                      285

Leu Asn Val Lys Cys Gln Leu Leu Glu Lys Glu Lys Glu Asp His Val  
                                  290                      295                      300

Asn Arg Asn Arg Glu His Asn Glu Asn Leu Asn Ala Glu Met Gln Asn  
                                  305                      310                      315                      320

Leu Lys Gln Lys Phe Ile Leu Glu Gln Gln Glu Arg Glu Lys Leu Gln  
                                  325                      330                      335

Gln Lys Glu Leu Gln Ile Asp Ser Leu Leu Gln Gln Glu Lys Glu Leu  
                                  340                      345                      350

Ser Ser Ser Leu His Gln Lys Leu Cys Ser Phe Gln Glu Glu Met Val  
                                  355                      360                      365

Lys Glu Lys Asn Leu Phe Glu Glu Glu Leu Lys Gln Thr Leu Asp Glu  
                                  370                      375                      380

Leu Asp Lys Leu Gln Gln Lys Glu Glu Gln Ala Glu Arg Leu Val Lys  
                                  385                      390                      395                      400

Gln Leu Glu Glu Glu Ala Lys Ser Arg Ala Glu Glu Leu Lys Leu Leu  
                                  405                      410                      415

Glu Glu Lys Leu Lys Gly Lys Glu Ala Glu Leu Glu Lys Ser Ser Ala  
                                  420                      425                      430

Ala His Thr Gln Ala Thr Leu Leu Leu Gln Glu Lys Tyr Asp Ser Met  
                                  435                      440                      445

Val Gln Ser Leu Glu Asp Val Thr Ala Gln Phe Glu Ser Tyr Lys Ala

450	455	460
Leu Thr Ala Ser Glu Ile Glu Asp Leu Lys Leu Glu Asn Ser Ser Leu		
465	470	475 480
Gln Glu Lys Ala Ala Lys Ala Gly Lys Asn Ala Glu Asp Val Gln His		
	485	490 495
Gln Ile Leu Ala Thr Glu Ser Ser Asn Gln Glu Tyr Val Arg Met Leu		
	500	505 510
Leu Asp Leu Gln Thr Lys Ser Ala Leu Lys Glu Thr Glu Ile Lys Glu		
	515	520 525
Ile Thr Val Ser Phe Leu Gln Lys Ile Thr Asp Leu Gln Asn Gln Leu		
	530	535 540
Lys Gln Gln Glu Glu Asp Phe Arg Lys Gln Leu Glu Asp Glu Glu Gly		
	545	550 555 560
Arg Lys Ala Glu Lys Glu Asn Thr Thr Ala Glu Leu Thr Glu Glu Ile		
	565	570 575
Asn Lys Trp Arg Leu Leu Tyr Glu Glu Leu Tyr Asn Lys Thr Lys Pro		
	580	585 590
Phe Gln Leu Gln Leu Asp Ala Phe Glu Val Glu Lys Gln Ala Leu Leu		
	595	600 605
Asn Glu His Gly Ala Ala Gln Glu Gln Leu Asn Lys Ile Arg Asp Ser		
	610	615 620
Tyr Ala Lys Leu Leu Gly His Gln Asn Leu Lys Gln Lys Ile Lys His		
	625	630 635 640
Val Val Lys Leu Lys Asp Glu Asn Ser Gln Leu Lys Ser Glu Val Ser		
	645	650 655
Lys Leu Arg Cys Gln Leu Ala Lys Lys Lys Gln Ser Glu Thr Lys Leu		
	660	665 670
Gln Glu Glu Leu Asn Lys Val Leu Gly Ile Lys His Phe Asp Pro Ser		
	675	680 685
Lys Ala Phe His His Glu Ser Lys Glu Asn Phe Ala Leu Lys Thr Pro		
	690	695 700
Leu Lys Glu Gly Asn Thr Asn Cys Tyr Arg Ala Pro Met Glu Cys Gln		
	705	710 715 720
Glu Ser Trp Lys		

<210> 81  
 <211> 713  
 <212> PRT

ALL INFORMATION CONTAINED HEREIN IS UNCLASSIFIED

Met 1	Ser	Phe	Pro	Lys 5	Ala	Pro	Leu	Lys	Arg 10	Phe	Asn	Asp	Pro	Ser	Gly
Cys	Ala	Pro	Ser 20	Pro	Gly	Ala	Tyr	Asp 25	Val	Lys	Thr	Ser	Glu 30	Ser	Thr
Lys	Gly	Pro 35	Val	Ser	Phe	Gln	Lys 40	Ser	Gln	Arg	Phe	Lys 45	Asn	Gln	Arg
Glu	Ser 50	Gln	Gln	Asn	Leu	Asn 55	Ile	Asp	Lys	Asp	Thr 60	Thr	Leu	Leu	Ala
Ser 65	Ala	Lys	Lys	Ala	Lys 70	Thr	Leu	Val	Ser	Lys 75	Lys	Glu	Ser	Gln	Lys 80
Asn	Asp	Lys	Asp	Val 85	Lys	Arg	Leu	Glu	Lys 90	Glu	Ile	His	Val	Leu 95	Leu
Gln	Glu	Arg	Gly 100	Thr	Gln	Asp	Lys	Arg 105	Ile	Gln	Asp	Met	Glu 110	Ser	Glu
Leu	Glu 115	Asn	Thr	Glu	Ala	Asn	Leu 120	Asn	Ala	Pro	Val	Thr 125	Glu	Lys	Pro
Ser 130	Leu	Ser	Ala	Asn	Asn	Ala 135	Ser	Leu	Glu	Lys	Arg 140	Leu	Thr	Glu	Leu
Thr 145	Arg	Ala	Asn	Glu	Leu 150	Leu	Lys	Ser	Lys	Phe 155	Ser	Glu	Asp	Ala	His 160
Gln	Lys	Asn	Met	Arg 165	Thr	Leu	Ser	Leu	Glu 170	Leu	Met	Lys	Leu	Arg 175	Asn
Lys	Arg	Glu	Thr 180	Lys	Met	Arg	Ser	Met 185	Met	Ala	Lys	Gln	Glu 190	Gly	Met
Glu	Leu 195	Lys	Leu	Gln	Ala	Thr	Gln 200	Lys	Asp	Leu	Ile	Glu 205	Ser	Lys	Gly
Lys 210	Ile	Val	Gln	Leu	Glu	Gly 215	Lys	Leu	Val	Ser	Ile 220	Glu	Lys	Glu	Lys
Ile 225	Asp	Glu	Lys	Ser	Glu 230	Thr	Glu	Lys	Leu	Leu 235	Glu	Tyr	Ile	Glu	Glu 240
Ile	Ser	Cys	Ala	Ser 245	Asp	Gln	Val	Glu	Lys 250	Tyr	Lys	Leu	Asp	Ile 255	Ala
Gln	Leu	Glu	Glu 260	Asp	Leu	Lys	Glu	Lys 265	Asp	Arg	Glu	Ile	Leu 270	Cys	Leu
Lys	Gln 275	Ser	Leu	Glu	Glu	Lys	Val 280	Ser	Phe	Ser	Lys	Gln 285	Ile	Glu	Asp



Leu Thr Val Lys Cys Gln Leu Leu Glu Ala Glu Arg Asp Asp Leu Val  
 290 295 300  
 Ser Lys Asp Arg Glu Arg Ala Glu Ser Leu Ser Ala Glu Met Gln Val  
 305 310 315 320  
 Leu Thr Glu Lys Leu Leu Leu Glu Arg Gln Glu Tyr Glu Lys Leu Gln  
 325 330 335  
 Gln Asn Glu Leu Gln Ser Gln Ser Leu Leu Gln Gln Glu Lys Glu Leu  
 340 345 350  
 Ser Ala His Leu Gln Gln Gln Leu Cys Ser Phe Gln Glu Glu Met Thr  
 355 360 365  
 Ser Glu Arg Asn Val Phe Lys Glu Gln Leu Lys Leu Ala Leu Asp Glu  
 370 375 380  
 Leu Asp Ala Val Gln Gln Lys Lys Glu Gln Ser Glu Lys Leu Val Lys  
 385 390 395 400  
 Gln Leu Glu Glu Glu Thr Lys Ser Thr Ala Glu Gln Leu Arg Arg Leu  
 405 410 415  
 Asp Asp Leu Leu Arg Glu Lys Glu Ile Glu Leu Glu Lys Arg Thr Ala  
 420 425 430  
 Ala His Ala Gln Ala Thr Val Ile Ala Gln Glu Lys Tyr Ser Asp Thr  
 435 440 445  
 Ala Gln Thr Leu Arg Asp Val Thr Ala Gln Leu Glu Ser Tyr Lys Ser  
 450 455 460  
 Ser Thr Leu Lys Glu Ile Glu Asp Leu Lys Leu Glu Asn Leu Thr Leu  
 465 470 475 480  
 Gln Glu Lys Val Ala Met Ala Glu Lys Arg Val Glu Asp Val Gln Gln  
 485 490 495  
 Gln Ile Leu Thr Ala Glu Ser Thr Asn Gln Glu Tyr Ala Lys Val Val  
 500 505 510  
 Gln Asp Leu Gln Asn Ser Ser Thr Leu Lys Glu Ala Glu Ile Lys Glu  
 515 520 525  
 Ile Thr Ser Ser Tyr Leu Glu Lys Ile Thr Asp Leu Gln Asn Gln Leu  
 530 535 540  
 Arg Gln Gln Asn Glu Asp Phe Arg Lys Gln Leu Glu Glu Glu Gly Ala  
 545 550 555 560  
 Lys Met Thr Glu Lys Glu Thr Ala Val Thr Glu Leu Thr Met Glu Ile  
 565 570 575  
 Asn Lys Trp Arg Leu Leu Tyr Glu Glu Leu Phe Asp Lys Thr Lys Pro  
 580 585 590



Val Asp Phe Gln Lys Asn Pro Glu Lys Ser Arg Gln Glu Ile Asn Phe  
 145 150 155 160  
 Trp Val Glu Cys Gln Ser Gln Gly Lys Ile Lys Glu Leu Phe Ser Lys  
 165 170 175  
 Asp Ala Ile Asn Ala Glu Thr Val Leu Val Leu Val Asn Ala Val Tyr  
 180 185 190  
 Phe Lys Ala Lys Trp Glu Thr Tyr Phe Asp His Glu Asn Thr Val Asp  
 195 200 205  
 Ala Pro Phe Cys Leu Asn Ala Asn Glu Asn Lys Ser Val Lys Met Met  
 210 215 220  
 Thr Gln Lys Gly Leu Tyr Arg Ile Gly Phe Ile Glu Glu Val Lys Ala  
 225 230 235 240  
 Gln Ile Leu Glu Met Arg Tyr Thr Lys Gly Lys Leu Ser Met Phe Val  
 245 250 255  
 Leu Leu Pro Ser His Ser Lys Asp Asn Leu Lys Gly Leu Glu Glu Leu  
 260 265 270  
 Glu Arg Lys Ile Thr Tyr Glu Lys Met Val Ala Trp Ser Ser Ser Glu  
 275 280 285  
 Asn Met Ser Glu Glu Ser Val Val Leu Ser Phe Pro Arg Phe Thr Leu  
 290 295 300  
 Glu Asp Ser Tyr Asp Leu Asn Ser Ile Leu Gln Asp Met Gly Ile Thr  
 305 310 315 320  
 Asp Ile Phe Asp Glu Thr Arg Ala Asp Leu Thr Gly Ile Ser Pro Ser  
 325 330 335  
 Pro Asn Leu Tyr Leu Ser Lys Ile Ile His Lys Thr Phe Val Glu Val  
 340 345 350  
 Asp Glu Asn Gly Thr Gln Ala Ala Ala Thr Gly Ala Val Val Ser  
 355 360 365  
 Glu Arg Ser Leu Arg Ser Trp Val Glu Phe Asn Ala Asn His Pro Phe  
 370 375 380  
 Leu Phe Phe Ile Arg His Asn Lys Thr Gln Thr Ile Leu Phe Tyr Gly  
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 Arg Val Cys Ser Pro  
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<210> 83  
 <211> 423  
 <212> PRT  
 <213> Mus musculus

<400> 83

Met Asp Ser Leu Thr Ala Ala Asn Asn Lys Phe Cys Phe Asp Phe Phe  
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 Arg Glu Ile Ser Lys Asp Asp Ala His Lys Asn Ile Phe Val Cys Pro  
 20 25 30  
 Leu Ser Leu Ser Ala Ala Phe Gly Met Val Arg Leu Gly Ala Arg Gly  
 35 40 45  
 Asp Ser Ala His Gln Ile Asp Glu Ala Leu His Phe Asn Glu Leu Ser  
 50 55 60  
 Lys Asp Glu His Lys Glu Pro Asn Asp Pro Ser Pro Gln Ser Glu Ser  
 65 70 75 80  
 Lys Ala Ser Asp Ser Ser Leu Glu Gly Gln Lys Gln Thr Ser Ala Ser  
 85 90 95  
 Gln Asp Gln Gln Gly Glu Ser Thr Asn Asp His Gln Leu Leu Gly Cys  
 100 105 110  
 His Phe Gly Lys Leu Leu Ser Arg Ile Asp Arg Asp Lys Ser Tyr Tyr  
 115 120 125  
 Thr Leu Ser Met Ala Asn Arg Leu Tyr Gly Glu Gln Glu Phe Pro Ile  
 130 135 140  
 Cys Ser Glu Tyr Ser Asp Asp Val Thr Glu Phe Phe His Thr Thr Val  
 145 150 155 160  
 Glu Ser Val Asp Phe Gln Lys Asp Ser Glu Lys Ser Arg Gln Glu Ile  
 165 170 175  
 Asn Phe Trp Val Glu Ser Gln Ser Gln Gly Lys Ile Lys Glu Leu Phe  
 180 185 190  
 Gly Lys Glu Ala Ile Asp Asn Ser Thr Val Leu Val Leu Val Asn Ala  
 195 200 205  
 Val Tyr Phe Lys Ala Lys Trp Glu Arg Glu Phe Asn Ser Glu Asn Thr  
 210 215 220  
 Val Asp Ala Ser Phe Cys Leu Asn Glu Asn Glu Lys Lys Thr Val Lys  
 225 230 235 240  
 Met Met Asn Gln Lys Gly Lys Phe Arg Ile Gly Phe Ile Asp Glu Leu  
 245 250 255  
 Gln Ala Gln Ile Leu Glu Met Lys Tyr Ala Met Gly Lys Leu Ser Met  
 260 265 270  
 Leu Val Leu Leu Pro Ser Cys Ser Glu Asp Asn Val Asn Ser Leu Gln  
 275 280 285  
 Glu Leu Glu Lys Lys Ile Asn His Glu Lys Leu Leu Ala Trp Ser Ser



Glu Gln Ser Arg Lys His Ile Asn Thr Trp Val Ala Glu Lys Thr Glu  
130 135 140

Gly Lys Ile Arg Asp Leu Leu Pro Ala Asn Ser Val Asn Pro Met Thr  
145 150 155 160

Arg Leu Val Leu Val Asn Ala Ile Tyr Phe Lys Gly Asn Trp Asp Thr  
165 170 175

Gln Phe Asn Lys Glu His Thr Glu Glu Arg Pro Phe Arg Val Ser Lys  
180 185 190

Asn Val Glu Lys Pro Val Gln Met Met Phe Lys Lys Ser Thr Cys Lys  
195 200 205

Ile Thr Tyr Ile Gly Glu Ile Ser Thr Gln Ile Leu Val Leu Pro Tyr  
210 215 220

Val Gly Gln Glu Leu Asn Met Val Ile Leu Leu Pro Ser Glu Ser Thr  
225 230 235 240

Asp Leu Asn Thr Val Glu Lys Ala Leu Thr Tyr Glu Lys Phe Ile Ala  
245 250 255

Trp Thr Lys Pro Asp Val Met Asp Glu Glu Glu Val Glu Val Phe Leu  
260 265 270

Pro Arg Phe Thr Leu Glu Glu Ser Tyr Asp Met Glu Glu Phe Leu Gln  
275 280 285

Glu Leu Gly Met Thr Asp Ala Phe Glu Glu Thr Arg Ala Asp Phe Ser  
290 295 300

Gly Met Ser Ser Gly Arg Gly Leu His Leu Ser Lys Val Met His Lys  
305 310 315 320

Ser Phe Val Glu Val Thr Glu Glu Gly Thr Glu Ala Ala Ala Ala Thr  
325 330 335

Gly Ala Val Val Met Met Arg Cys Leu Met Val Val Pro Arg Phe Asn  
340 345 350

Ala Asn His Pro Phe Leu Phe Phe Ile Gln His Ser Lys Thr Gly Ala  
355 360 365

Ile Leu Phe Cys Gly Arg Phe Cys Ser Pro  
370 375

<210> 85

<211> 379

<212> PRT

<213> Mus musculus

<400> 85

Met Glu Gln Leu Ser Ser Ala Asn Thr Leu Phe Ala Leu Glu Leu Phe  
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Gln Thr Leu Asn Glu Ser Ser Pro Thr Gly Asn Ile Phe Phe Ser Pro  
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 Phe Ser Ile Ser Ser Ala Leu Ala Met Val Ile Leu Gly Ala Lys Gly  
 35 40 45  
 Ser Thr Ala Ala Gln Leu Ser Lys Thr Phe His Phe Asp Ser Val Glu  
 50 55 60  
 Asp Ile His Ser Arg Phe Gln Ser Leu Asn Ala Glu Val Ser Lys Arg  
 65 70 75 80  
 Gly Ala Ser His Thr Leu Lys Leu Ala Asn Arg Leu Tyr Gly Glu Lys  
 85 90 95  
 Thr Tyr Asn Phe Leu Pro Glu Tyr Leu Ala Ser Thr Gln Lys Met Tyr  
 100 105 110  
 Gly Ala Asp Leu Ala Pro Val Asp Phe Leu His Ala Ser Glu Asp Ala  
 115 120 125  
 Arg Lys Glu Ile Asn Gln Trp Val Lys Gly Gln Thr Glu Gly Lys Ile  
 130 135 140  
 Pro Glu Leu Leu Ser Val Gly Val Val Asp Ser Met Thr Lys Leu Val  
 145 150 155 160  
 Leu Val Asn Ala Ile Tyr Phe Lys Gly Met Trp Glu Glu Lys Phe Met  
 165 170 175  
 Thr Glu Asp Thr Thr Asp Ala Pro Phe Arg Leu Ser Lys Lys Asp Thr  
 180 185 190  
 Lys Thr Val Lys Met Met Tyr Gln Lys Lys Lys Phe Pro Phe Gly Tyr  
 195 200 205  
 Ile Ser Asp Leu Lys Cys Lys Val Leu Glu Met Pro Tyr Gln Gly Gly  
 210 215 220  
 Glu Leu Ser Met Val Ile Leu Leu Pro Lys Asp Ile Glu Asp Glu Ser  
 225 230 235 240  
 Thr Gly Leu Lys Lys Ile Glu Lys Gln Ile Thr Leu Glu Lys Leu Leu  
 245 250 255  
 Glu Trp Thr Lys Arg Glu Asn Leu Glu Phe Ile Asp Val His Val Lys  
 260 265 270  
 Leu Pro Arg Phe Lys Ile Glu Glu Ser Tyr Thr Leu Asn Ser Asn Leu  
 275 280 285  
 Gly Arg Leu Gly Val Gln Asp Leu Phe Ser Ser Ser Lys Ala Asp Leu  
 290 295 300  
 Ser Gly Met Ser Gly Ser Arg Asp Leu Phe Ile Ser Lys Ile Val His  
 305 310 315 320

Lys Ser Phe Val Glu Val Asn Glu Glu Gly Thr Glu Ala Ala Ala Ala  
325 330 335

Thr Gly Gly Ile Ala Thr Phe Cys Met Leu Leu Pro Glu Glu Glu Phe  
340 345 350

Thr Val Asp His Pro Phe Ile Phe Phe Ile Arg His Asn Pro Thr Ser  
355 360 365

Asn Val Leu Phe Leu Gly Arg Val Cys Ser Pro  
370 375

<210> 86

<211> 379

<212> PRT

<213> Mus musculus

<220>

<221> VARIANT

<222> (204)

<223> Wherein Xaa is any amino acid as defined in the  
specification.

<400> 86

Met Glu Gln Leu Ser Ser Ala Asn Thr Leu Phe Ala Leu Glu Leu Phe  
1 5 10 15

Gln Thr Leu Asn Glu Ser Ser Pro Thr Gly Asn Ile Phe Phe Ser Pro  
20 25 30

Phe Ser Ile Ser Ser Ala Leu Ala Met Val Ile Leu Gly Ala Lys Gly  
35 40 45

Ser Thr Ala Ala Gln Leu Ser Lys Thr Phe His Phe Asp Ser Val Glu  
50 55 60

Asp Ile His Ser Arg Phe Gln Ser Gln Asn Ala Glu Val Ser Lys Arg  
65 70 75 80

Gly Ala Ser His Thr Leu Lys Leu Ala Asn Arg Leu Tyr Gly Glu Lys  
85 90 95

Thr Tyr Asn Phe Leu Pro Glu Tyr Leu Ala Ser Thr Gln Lys Met Tyr  
100 105 110

Gly Ala Asp Leu Ala Pro Val Asp Phe Leu His Ala Ser Glu Asp Ala  
115 120 125

Arg Lys Glu Ile Asn Gln Trp Val Lys Gly Gln Thr Glu Gly Lys Ile  
130 135 140

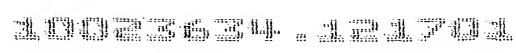
Pro Glu Leu Leu Ser Val Gly Val Val Asp Ser Met Thr Lys Leu Val  
145 150 155 160

Leu Val Asn Ala Ile Tyr Phe Lys Gly Met Trp Glu Glu Lys Phe Met









Leu Ile Tyr Asp Asp Pro Thr Gly Ser Ile Leu Phe Met Gly Lys Val  
355 360 365

Val Asn Pro  
370

<210> 88  
<211> 360  
<212> PRT  
<213> Homo sapiens

<400> 88  
Phe Asp Leu Tyr Lys Glu Leu Ala Lys Glu Ser Pro Asp Lys Asn Ile  
1 5 10 15

Phe Phe Ser Pro Val Ser Ile Ser Ser Ala Leu Ala Met Leu Ser Leu  
20 25 30

Gly Ala Lys Gly Ser Thr Ala Thr Gln Ile Leu Glu Val Leu Gly Phe  
35 40 45

Asn Leu Thr Glu Thr Ser Glu Ala Asp Ile His Gln Gly Phe Gln His  
50 55 60

Leu Leu His Leu Leu Asn Arg Pro Asp Asn Lys Leu Gln Leu Lys Thr  
65 70 75 80

Ala Asn Ala Leu Phe Val Asp Lys Ser Leu Lys Leu Leu Asp Ser Phe  
85 90 95

Leu Glu Asp Val Lys Lys Leu Tyr Gly Ala Glu Val Gln Ser Val Asp  
100 105 110

Phe Ser Asp Pro Ala Glu Glu Ala Lys Lys Gln Ile Asn Asp Trp Val  
115 120 125

Lys Lys Lys Thr Gln Gly Lys Ile Lys Asp Leu Leu Ser Asp Leu Asp  
130 135 140

Pro Asp Thr Arg Leu Val Leu Val Asn Ala Ile Tyr Phe Lys Gly Lys  
145 150 155 160

Trp Lys Thr Pro Phe Asp Pro Glu Asn Thr Arg Glu Glu Asp Phe Tyr  
165 170 175

Val Asp Glu Thr Thr Thr Val Lys Val Pro Met Met Ser Gln Thr Gly  
180 185 190

Arg Thr Phe Arg Tyr Gly Arg Asp Glu Glu Leu Asn Cys Gln Val Leu  
195 200 205

Glu Leu Pro Tyr Lys Gly Asn Ala Ser Met Leu Ile Ile Leu Pro Asp  
210 215 220

Glu Gly Gly Leu Glu Thr Val Glu Lys Ala Leu Thr Pro Glu Thr Leu  
225 230 235 240



130	135	140
Thr Ser Ile Glu Val	Val Ala Ala Asp Thr	Pro Ala Pro Phe Ser Arg
145	150	155 160
Tyr Gln Ala Gln Asn Phe Thr Leu Val Cys Ile Val Ser Gly Gly Lys		
	165	170 175
Pro Ala Pro Met Val Tyr Phe Lys Arg Asp Gly Glu Pro Ile Asp Ala		
	180	185 190
Val Pro Leu Ser Glu Pro Pro Ala Ala Ser Ser Gly Pro Leu Gln Asp		
	195	200 205
Ser Arg Pro Phe Arg Ser Leu Leu His Arg Asp Leu Asp Asp Thr Lys		
	210	215 220
Met Gln Lys Ser Leu Ser Leu Leu Asp Ala Glu Asn Arg Gly Gly Arg		
	225	230 235 240
Pro Tyr Thr Glu Arg Pro Ser Arg Gly Leu Thr Pro Asp Pro Asn Ile		
	245	250 255
Leu Leu Gln Pro Thr Thr Glu Asn Ile Pro Glu Thr Val Val Ser Arg		
	260	265 270
Glu Phe Pro Arg Trp Val His Ser Ala Glu Pro Thr Tyr Phe Leu Arg		
	275	280 285
His Ser Arg Thr Pro Ser Ser Asp Gly Thr Val Glu Val Arg Ala Leu		
	290	295 300
Leu Thr Trp Thr Leu Asn Pro Gln Ile Asp Asn Glu Ala Leu Phe Ser		
	305	310 315 320
Cys Glu Val Lys His Pro Ala Leu Ser Met Pro Met Gln Ala Glu Val		
	325	330 335
Thr Leu Val Ala Pro Lys Gly Pro Lys Ile Val Met Thr Pro Ser Arg		
	340	345 350
Ala Arg Val Gly Asp Thr Val Arg Ile Leu Val His Gly Phe Gln Asn		
	355	360 365
Glu Val Phe Pro Glu Pro Met Phe Thr Trp Thr Arg Val Gly Ser Arg		
	370	375 380
Leu Leu Asp Gly Ser Ala Glu Phe Asp Gly Lys Glu Leu Val Leu Glu		
	385	390 395 400
Arg Val Pro Ala Glu Leu Asn Gly Ser Met Tyr Arg Cys Thr Ala Gln		
	405	410 415
Asn Pro Leu Gly Ser Thr Asp Thr His Thr Arg Leu Ile Val Phe Glu		
	420	425 430
Asn Pro Asn Ile Pro Arg Gly Thr Glu Asp Ser Asn Gly Ser Ile Gly		

435 440 445

Pro Thr Gly Ala Arg Leu Thr Leu Val Leu Ala Leu Thr Val Ile Leu  
450 455 460

Glu Leu Thr  
465

<210> 90  
<211> 404  
<212> PRT  
<213> *Macaca fascicularis*

<400> 90  
Met Arg Ala Ala Pro Ser Leu Arg Arg Cys Val Cys Leu Leu Leu Ala  
1 5 10 15

Ala Ile Leu Asp Leu Ala Cys Gly Tyr Leu Thr Val Asn Ile Glu Pro  
20 25 30

Leu Pro Pro Val Val Ala Gly Asp Ala Val Thr Leu Lys Cys Asn Phe  
35 40 45

Lys Thr Asp Gly Arg Met Arg Glu Ile Val Trp Tyr Arg Val Thr Asp  
50 55 60

Gly Gly Thr Ile Lys Gln Lys Ile Phe Thr Phe Asp Ala Met Phe Ser  
65 70 75 80

Thr Asn Tyr Ser His Met Glu Asn Tyr Arg Lys Arg Glu Asp Leu Val  
85 90 95

Tyr Gln Ser Thr Val Arg Leu Pro Glu Val Arg Ile Ser Asp Asn Gly  
100 105 110

Pro Tyr Glu Cys His Val Gly Ile Tyr Asp Arg Ala Thr Arg Glu Lys  
115 120 125

Val Val Leu Ala Ser Gly Asn Ile Phe Leu Asn Val Met Ala Pro Pro  
130 135 140

Thr Ser Ile Glu Val Val Ala Ala Asp Thr Pro Ala Pro Phe Ser Arg  
145 150 155 160

Tyr Gln Ala Gln Asn Phe Thr Leu Val Cys Ile Val Ser Gly Gly Lys  
165 170 175

Pro Ala Pro Met Val Tyr Phe Lys Arg Asp Gly Glu Pro Ile Asp Ala  
180 185 190

Val Pro Leu Ser Glu Pro Pro Ala Ala Ser Ser Gly Pro Leu Gln Asp  
195 200 205

Ser Arg Pro Phe Arg Ser Leu Leu His Arg Asp Leu Asp Asp Thr Lys  
210 215 220

Met Gln Lys Ser Leu Ser Leu Leu Asp Ala Glu Asn Arg Gly Gly Arg  
 225 230 235 240

Pro Tyr Thr Glu Arg Pro Ser His Gly Leu Thr Pro Asp Pro Asn Ile  
 245 250 255

Leu Leu Gln Pro Thr Thr Glu Asn Ile Pro Glu Thr Val Val Ser Arg  
 260 265 270

Glu Phe Pro Arg Trp Val His Ser Ala Glu Pro Thr Tyr Phe Leu Arg  
 275 280 285

His Ser Arg Thr Pro Ser Ser Asp Gly Thr Val Glu Val Arg Ala Leu  
 290 295 300

Leu Thr Trp Thr Leu Asn Pro Gln Ile Asp Asn Glu Ala Leu Phe Ser  
 305 310 315 320

Cys Glu Val Lys His Pro Ala Leu Ser Met Pro Met Gln Ala Glu Val  
 325 330 335

Thr Leu Val Ala Pro Lys Gly Pro Lys Ile Val Met Thr Pro Ser Arg  
 340 345 350

Ala Arg Val Gly Asp Thr Val Arg Ile Leu Val His Gly Phe Gln Asn  
 355 360 365

Glu Val Phe Pro Glu Pro Met Phe Thr Trp Thr Arg Val Gly Ser Arg  
 370 375 380

Leu Leu Asp Gly Ser Ala Glu Phe Asp Gly Lys Glu Leu Val Leu Glu  
 385 390 395 400

Arg Val Pro Ala

<210> 91  
 <211> 80  
 <212> PRT  
 <213> Homo sapiens

<400> 91  
 Val Thr Asp Gly Gly Thr Ile Lys Gln Lys Ile Phe Thr Phe Asp Ala  
 1 5 10 15

Met Phe Ser Thr Asn Tyr Ser His Met Glu Asn Tyr Arg Lys Arg Glu  
 20 25 30

Asp Leu Val Tyr Gln Ser Thr Val Arg Leu Pro Glu Val Arg Ile Ser  
 35 40 45

Asp Asn Gly Pro Tyr Glu Cys His Val Gly Ile Tyr Asp Arg Ala Thr  
 50 55 60

Arg Glu Lys Val Val Leu Ala Ser Gly Asn Ile Phe Leu Asn Val Met  
 65 70 75 80

Met Thr Thr Lys Ala Pro Thr Phe Thr Gln Pro Leu Gln Ser Val Val  
1 5 10 15

Pro Val Pro Glu Val Ser Trp Tyr Arg Asp Gly Gln Val Leu Ser Ala  
35 40 45

Leu Val Ile Pro Ser Val Thr Glu Ala Asn Ser Gly Arg Tyr Thr Ile  
65 70 75 80

Val Thr Ala Gly Thr Ala Pro Pro Asn Phe Ser Gln Arg Leu Gln Ser  
100 105 110

Gly Ile Pro Thr Pro Val Val Lys Phe Tyr Arg Asp Gly Val Glu Ile  
130 135 140

Leu Ile Ile Ala Glu Ala Tyr Pro Glu Asp Ser Gly Thr Tyr Ser Val  
165 170 175

Ile Gln Gly Glu Glu Glu Ala Val Pro Ala Lys Lys Thr Lys Thr Ile  
195 200 205

Lys Ile Glu Thr His Phe Asp Ala Arg Ser Leu Thr Ser Val Glu Met  
225 230 235 240

132





545                      550                      555                      560

Ser Pro His Phe Thr Val Ser Lys Ile Ala Val Pro Lys Pro Asp His  
565                      570                      575

Thr Tyr Glu Val Ser Ile Ala Gly Ser Ala Met Ala Thr Leu Glu Lys  
580                      585                      590

Glu Leu Ser Ala Thr Ser Ala Ala Gln Lys Ile Thr Lys Pro Val Lys  
595                      600                      605

Pro Pro Gln Leu Lys Pro His Glu Val Lys Ile Lys Pro Glu Ser Ala  
610                      615                      620

Pro Pro Gln Phe Pro Phe Thr Glu Ala Ala Glu Thr Tyr Lys Ala His  
625                      630                      635                      640

Tyr Asp Val Glu Thr Lys Lys Glu Val Asp Val Ser Ile Lys Gly Glu  
645                      650                      655

Ala Val Arg Glu Asp His Leu Leu Leu Arg Lys Glu Ser Glu Ala Lys  
660                      665                      670

Val Thr Glu Thr Ala Arg Val Pro Val Pro Ala Glu Ile Pro Val Thr  
675                      680                      685

Pro Pro Thr Leu Val Trp Gly Leu Lys Asn Lys Thr Val Thr Glu Gly  
690                      695                      700

Glu Ser Val Thr Leu Glu Cys His Ile Ser Gly His Pro Gln Pro Thr  
705                      710                      715                      720

Val Thr Trp Tyr Arg Glu Asp Tyr Lys Ile Glu Ser Ser Met Asp Phe  
725                      730                      735

Gln Ile Thr Phe Lys Ala Gly Leu Ala Arg Leu Val Ile Arg Glu Ala  
740                      745                      750

Phe Ala Glu Asp Ser Gly Arg Phe Thr Cys Thr Ala Thr Asn Lys Ala  
755                      760                      765

Gly Ser Val Ser Thr Ser Cys His Leu His Val Lys Val Ser Glu Glu  
770                      775                      780

Thr Glu Thr Arg Glu Thr Ile Ser Glu Lys Val Val Thr Glu Glu Lys  
785                      790                      795                      800

Ser Tyr Val Glu Thr Lys Asp Val Val Met Glu Asp Val Ser Ala Ala  
805                      810                      815

Ala Glu Glu Val Ser Gly Glu Pro Val Pro Pro Phe Phe Ile Arg Lys  
820                      825                      830

Pro Val Val His Lys Leu Ile Glu Gly Gly Ser Ile Ile Phe Glu Cys  
835                      840                      845

Gln Val Gly Gly Asn Pro Lys Pro His Val Leu Trp Lys Lys Gly Gly

850					855					860					
Val	Pro	Leu	Thr	Thr	Gly	Tyr	Arg	Tyr	Lys	Val	Ser	Tyr	Lys	Arg	Glu
865					870					875					880
Thr	Gly	Glu	Cys	Lys	Leu	Glu	Ile	Ser	Met	Thr	Phe	Ala	Asp	Asp	Ala
				885					890					895	
Gly	Glu	Tyr	Thr	Ile	Val	Ile	Arg	Asn	Lys	Phe	Gly	Glu	Ala	Ser	Ala
			900					905					910		
Thr	Val	Ser	Leu	Leu	Glu	Glu	Ala	Asp	Tyr	Glu	Ala	Tyr	Ile	Lys	Ser
		915					920					925			
Gln	Gln	Glu	Met	Met	Tyr	Gln	Thr	Gln	Val	Thr	Ala	Tyr	Val	Gln	Glu
	930					935					940				
Pro	Lys	Val	Ala	Glu	Val	Ala	Pro	Pro	Ile	Ser	Tyr	Gly	Asp	Phe	Asp
945					950					955					960
Lys	Glu	Tyr	Glu	Lys	Glu	Gln	Ala	Leu	Ile	Arg	Lys	Lys	Met	Ala	Lys
				965					970					975	
Asp	Thr	Val	Met	Val	Arg	Thr	Phe	Val	Glu	Asp	Glu	Glu	Phe	His	Ile
			980					985					990		
Ser	Ser	Phe	Glu	Glu	Arg	Leu	Ile	Lys	Glu	Ile	Glu	Leu	Arg	Ile	Ile
		995				1000						1005			
Lys	Thr	Thr	Leu	Asp	Glu	Leu	Leu	Glu	Glu	Asp	Gly	Glu	Glu	Met	Met
	1010					1015					1020				
Ile	Asp	Ile	Ser	Glu	Ser	Glu	Ala	Ile	Gly	Ala	Gly	Phe	Asp	Leu	Arg
1025				1030						1035					1040
Leu	Lys	Asn	Tyr	Arg	Thr	Phe	Glu	Gly	Thr	Gly	Val	Thr	Phe	His	Cys
				1045				1050						1055	
Lys	Thr	Thr	Gly	Tyr	Pro	Leu	Pro	Lys	Ile	Ala	Trp	Tyr	Lys	Asp	Gly
		1060					1065						1070		
Lys	Arg	Ile	Arg	His	Gly	Glu	Arg	Tyr	His	Met	Glu	Val	Leu	Gln	Asp
	1075					1080					1085				
Gly	Ser	Ala	Ser	Leu	Arg	Leu	Pro	Val	Val	Leu	Pro	Glu	Asp	Glu	Gly
	1090				1095					1100					
Ile	Tyr	Thr	Val	Phe	Ala	Ser	Asn	Met	Lys	Gly	Asn	Ala	Ile	Cys	Ser
1105				1110						1115				1120	
Ala	Lys	Leu	Tyr	Val	Glu	Pro	Val	Ala	Pro	Thr	Ala	Thr	Pro	Gly	Tyr
				1125				1130					1135		
Met	Pro	Gly	Pro	Glu	Val	Met	Arg	Arg	Tyr	Arg	Ser	Ile	Ser	Pro	Arg
		1140					1145					1150			
Ser	Pro	Ser	Arg	Ser	Pro	Ala	Arg	Ser	Ser	Pro	Ser	Cys	Ser	Pro	Ala



1460                      1465                      1470  
 Leu Glu Ala Ala Asn Arg Leu Arg Met Ile Asn Glu Phe Gly Tyr Cys  
       1475                      1480                      1485  
 Ser Leu Asp Tyr Gly Val Ala Tyr Ser Arg Asp Ser Gly Val Ile Thr  
       1490                      1495                      1500  
 Cys Arg Ala Thr Asn Lys Tyr Gly Thr Asp His Thr Ser Ala Thr Leu  
       1505                      1510                      1515                      1520  
 Ile Val Lys Asp Glu Lys Ser Leu Val Glu Glu Ser Gln Leu Pro Glu  
               1525                      1530                      1535  
 Gly Arg Arg Gly Met Gln Arg Ile Glu Glu Leu Glu Arg Met Ala His  
               1540                      1545                      1550  
 Glu Gly Ala Leu Pro Ala Val Ala Val Asp Gln Lys Glu Lys Gln Lys  
               1555                      1560                      1565  
 Pro Glu Leu Val Leu Val Pro Glu Pro Ala Arg Val Leu Glu Gly Glu  
               1570                      1575                      1580  
 Thr Ala Arg Phe Arg Cys Arg Val Thr Gly Tyr Pro Leu Pro Lys Val  
       1585                      1590                      1595                      1600  
 Asn Trp Tyr Leu Asn Ser Gln Leu Ile Arg Lys Ser Lys Arg Phe Arg  
               1605                      1610                      1615  
 Leu Arg Tyr Asp Gly Ile His Tyr Leu Asp Ile Val Asp Cys Lys Ser  
               1620                      1625                      1630  
 Tyr Asp Thr Gly Glu Val Lys Val Thr Ala Glu Asn Pro Glu Gly Phe  
               1635                      1640                      1645  
 Ile Glu His Lys Val Lys Leu Glu Ile Gln Gln Arg Glu Asp Phe Arg  
               1650                      1655                      1660  
 Ser Val Leu Arg Arg Ala Pro Glu Pro Arg His Glu Pro Val Val Thr  
       1665                      1670                      1675                      1680  
 Glu Pro Gly Lys Leu Leu Phe Glu Val Gln Lys Ile Asp Lys Pro Ala  
               1685                      1690                      1695  
 Glu Ala Thr Thr Lys Glu Val Val Lys Leu Lys Arg Ala Glu Arg Ile  
               1700                      1705                      1710  
 Thr His Glu Lys Leu Ser Glu Glu Ser Glu Glu Leu Arg Ser Lys Phe  
               1715                      1720                      1725  
 Lys Arg Arg Thr Glu Glu Gly Tyr Tyr Glu Ala Ile Thr Ala Val Glu  
               1730                      1735                      1740  
 Leu Lys Ser Arg Lys Lys Asp Glu Ser Tyr Glu Glu Met Leu Lys Lys  
       1745                      1750                      1755                      1760  
 Thr Lys Glu Glu Leu Leu His Trp Thr Lys Glu Ile Pro Glu Glu Glu

Lys Lys Ala Leu Pro Pro Glu Gly Lys Ile Thr Ile Pro Thr Phe Lys  
 1780 1785 1790  
 Pro Glu Lys Val Glu Leu Ser Pro Ser Met Glu Ala Pro Lys Ile Phe  
 1795 1800 1805  
 Glu Arg Ile Gln Ser Gln Thr Val Ala Gln Gly Thr Asp Ala His Phe  
 1810 1815 1820  
 Arg Val Arg Val Val Gly Lys Pro Asp Pro Glu Cys Gln Trp Phe Arg  
 1825 1830 1835 1840  
 Asn Gly Val Gln Ile Glu Arg Thr Asp Arg Ile Tyr Trp Tyr Trp Pro  
 1845 1850 1855  
 Glu Asp Asn Val Cys Glu Leu Val Ile Arg Asp Val Thr Ala Asp Asp  
 1860 1865 1870  
 Ser Ala Ser Ile Met Val Lys Ala Val Asn Ile Ala Gly Glu Thr Ser  
 1875 1880 1885  
 Ser His Ala Phe Leu Leu Val Gln Ala Lys Gln Leu Ile Ser Phe Ile  
 1890 1895 1900  
 Gln Asn Leu Gln Asp Val Val Ala Lys Glu Arg Asp Ser Met Ala Thr  
 1905 1910 1915 1920  
 Phe Glu Cys Glu Thr Ser Glu Pro Phe Ile Lys Val Lys Trp Phe Lys  
 1925 1930 1935  
 Asn Gly Ile Glu Ile His Ser Gly Glu Lys Tyr Arg Met His Ser Asp  
 1940 1945 1950  
 Arg Lys Ala His Phe Leu Ser Val Leu Ala Val Glu Met Ser Asp Ala  
 1955 1960 1965  
 Asp Asp Tyr Ser Cys Ala Leu Val Glu Asp Glu Ser Val Lys Thr Thr  
 1970 1975 1980  
 Ala Lys Leu Ile Val Glu Gly Ala Val Val Glu Phe Ile Lys Glu Leu  
 1985 1990 1995 2000  
 Glu Asp Val Glu Val Pro Glu Ser Phe Thr Gly Glu Leu Glu Cys Glu  
 2005 2010 2015  
 Val Ser Pro Glu Asp Ile Glu Gly Lys Trp Tyr His Gly Asp Val Glu  
 2020 2025 2030  
 Leu Ser Ser Asn His Lys Tyr Val Leu Ala Ser Arg Arg Gly Arg Arg  
 2035 2040 2045  
 Ile Leu Thr Ile Lys Asp Val Asn Lys Asp Asp Gln Gly Glu Tyr Ser  
 2050 2055 2060  
 Phe Val Val Asp Gly Lys Arg Thr His Cys Lys Leu Lys Met Lys Pro

2065                      2070                      2075                      2080  
 Arg Pro Met Thr Ile Leu Gln Gly Leu Thr Asp Gln Lys Val Cys Glu  
                                  2085                      2090                      2095  
 Gly Asp Ile Val Gln Leu Glu Val Lys Val Ser Val Glu Asn Val Glu  
                                  2100                      2105                      2110  
 Gly Val Trp Met Lys Asp Gly His Glu Ile Gln Ser Ser Asp Arg Ile  
                                  2115                      2120                      2125  
 His Ile Val Leu Asp Lys Gln Ser His Met Leu Leu Ile Glu Asp Ala  
                                  2130                      2135                      2140  
 Thr Gln Glu Asp Ser Gly Thr Tyr Ser Phe Ser Ile Pro Gly Leu Glu  
 2145                      2150                      2155                      2160  
 Leu Ser Thr Thr Gly Gln Val Thr Val Tyr Ser Val Glu Ile Ile Val  
                                  2165                      2170                      2175  
 Pro Leu Lys Asp Val His Val Val Glu Gly Thr Lys Ala Ile Leu Glu  
                                  2180                      2185                      2190  
 Cys Lys Val Ser Ala Pro Asp Val Thr Ser Ser Lys Trp Tyr Leu Asn  
                                  2195                      2200                      2205  
 Asp His Gln Ile Lys Pro Asp Glu Arg Val Gln Ala Val Cys Lys Gly  
                                  2210                      2215                      2220  
 Thr Lys Gln Arg Leu Val Ile Thr Arg Thr His Ala Ser Asp Glu Gly  
 2225                      2230                      2235                      2240  
 His Tyr Lys Leu Met Val Gly Lys Val Glu Thr Ser Cys Asn Val Thr  
                                  2245                      2250                      2255  
 Val Glu Glu Ile Glu Ile Ile Arg Gly Leu His Asp Ile Thr Cys Thr  
                                  2260                      2265                      2270  
 Glu Thr Gln Asn Val Ser Phe Glu Val Glu Leu Ser His Ser Gly Ile  
                                  2275                      2280                      2285  
 Asp Val Ile Trp His Phe Lys Gly Gln Glu Ile Lys Ala Gly Pro Lys  
                                  2290                      2295                      2300  
 Tyr Lys Ile Glu Ala Arg Gly Lys Ile Tyr Lys Leu Thr Val Val Lys  
 2305                      2310                      2315                      2320  
 Met Met Lys Asp Asp Glu Gly Glu Tyr Val Phe Tyr Ala Gly Gly Lys  
                                  2325                      2330                      2335  
 Lys Thr Ser Gly Lys Leu Ile Val Ala Gly Gly Ala Ile Ser Lys Pro  
                                  2340                      2345                      2350  
 Leu Ala Asp Leu Thr Val Ala Glu Ser Gln Arg Ala Val Phe Glu Cys  
                                  2355                      2360                      2365  
 Glu Val Ala Asn Pro Glu Ser Glu Gly Gln Trp Leu Lys Asn Gly Lys

2370                      2375                      2380

Pro Leu Pro Met Thr Asp Gln Tyr Arg Ala Glu Thr Asp Gly Val Lys  
 2385                      2390                      2395                      2400

Arg Arg Leu Asn Ile Pro Ala Ala Lys Met Asp Asp Met Gly Glu Tyr  
                     2405                      2410                      2415

Ser Tyr Glu Ile Ala Ser Ser Lys Thr Ser Ala Lys Leu His Val Glu  
                     2420                      2425                      2430

Ala Val Lys Ile Lys Lys Thr Leu Lys Asn Leu Thr Val Thr Glu Thr  
                     2435                      2440                      2445

Gln Glu Ala Val Phe Ser Val Glu Leu Ser His Pro Asp Val Lys Gly  
                     2450                      2455                      2460

Ala Leu Trp Ile Lys Asn Gly Val Glu Leu Glu Ser Asn Asp Lys Tyr  
                     2465                      2470                      2475                      2480

Glu Ile Ser Val Lys Gly Thr Val His Thr Leu Lys Ile Lys His Cys  
                     2485                      2490                      2495

Val Val Thr Asp Glu Ser Val Tyr Ser Phe Lys Leu Gly Lys Ile Gly  
                     2500                      2505                      2510

Ala Asn Ala Arg Leu His Val Glu Thr Val Lys Ile Ile Lys Lys Pro  
                     2515                      2520                      2525

Lys Asp Val Thr Ala Leu Glu Asn Ala Val Val Ser Phe Glu Leu Ser  
                     2530                      2535                      2540

Val Ser His Asp Thr Val Pro Val Arg Trp Phe His Lys Asn Val Glu  
                     2545                      2550                      2555                      2560

Leu Lys Gln Ser Asp Lys Tyr Lys Met Ile Ser Gln Arg Lys Val His  
                     2565                      2570                      2575

Lys Leu Met Leu His Asn Ile Ser Pro Ala Asp Ala Gly Glu Tyr Thr  
                     2580                      2585                      2590

Ala Phe Val Gly Gln Leu Glu Cys Lys Ala Lys Leu Phe Val Glu Thr  
                     2595                      2600                      2605

Ile His Ile Thr Lys Thr Met Lys Ser Ile Glu Ile Pro Glu Thr Lys  
                     2610                      2615                      2620

Thr Ala Ser Phe Gln Cys Glu Val Ser His Phe Asn Val Pro Ser Val  
                     2625                      2630                      2635                      2640

Trp Leu Lys Asn Gly Val Glu Ile Glu Met Ser Glu Lys Phe Lys Ile  
                     2645                      2650                      2655

Val Val Gln Gly Lys Leu His Gln Leu Asn Ile Met Asn Thr Ser Ser  
                     2660                      2665                      2670

Glu Asp Ser Ala Glu Tyr Thr Phe Val Cys Gly Asn Asp Arg Val Ser



2675	2680	2685
Ala Thr Leu Thr Val Lys Pro Ile Leu Ile Thr Ser Met Leu Glu Asp 2690	2695	2700
Ile Asn Ala Glu Glu Lys Asp Thr Ile Thr Phe Glu Val Thr Val Asn 2705	2710	2715 2720
Tyr Glu Gly Ile Ser Tyr Lys Trp Leu Lys Asn Gly Val Glu Ile Lys 2725	2730	2735
Ser Thr Asp Lys Cys Gln Ile Arg Thr Lys Lys Leu Thr His Ser Leu 2740	2745	2750
Ser Ile Arg Asn Val His Phe Gly Asp Ala Ala Glu Tyr Ser Phe Val 2755	2760	2765
Ala Gly Lys Ala Ala Ser Ser Ala Thr Leu Tyr Val Glu Ala Arg His 2770	2775	2780
Ile Glu Phe Arg Lys His Ile Lys Asp Ile Lys Val Val Glu Lys Lys 2785	2790	2795 2800
Arg Ala Ile Phe Glu Cys Glu Ile Ser Glu Pro Asp Val Gln Val Gln 2805	2810	2815
Trp Met Lys Asp Gly Gln Glu Leu Gln Ile Gly Asp Arg Met Lys Ile 2820	2825	2830
Gln Arg Glu Lys Tyr Val His Arg Leu Ile Ile Pro Ser Thr Lys Met 2835	2840	2845
Ser Asp Ala Gly Gln Tyr Thr Val Val Ala Gly Gly Asn Thr Ser Ser 2850	2855	2860
Ala Asn Leu Ile Val Glu Gly Arg Asp Val Arg Ile Arg Ser Ile Arg 2865	2870	2875 2880
Lys Glu Ile Gln Val Ile Glu Arg Gln Arg Ala Glu Ile Glu Phe Glu 2885	2890	2895
Val Asn Glu Asp Asp Ile Glu Pro Gln Trp Tyr Lys Asp Gly Ile Glu 2900	2905	2910
Ile Asn Phe His Tyr Glu Glu Arg Tyr Ser Tyr Val Val Glu Arg Arg 2915	2920	2925
Ile His Arg Met Ser Ile Phe Glu Thr Thr Tyr Ser Asp Ala Gly Glu 2930	2935	2940
Tyr Thr Phe Val Ala Gly Arg Asn Arg Ser Ser Val Val Leu Tyr Val 2945	2950	2955 2960
Asn Ala Pro Glu Pro Pro Gln Ile Ile Gln Glu Leu Gln Pro Thr Thr 2965	2970	2975
Val Glu Ser Gly Lys Pro Ala Arg Phe Cys Ala Ile Ile Ser Gly Lys		

2980					2985					2990					
Pro	Gln	Pro	Lys	Val	Ser	Trp	Tyr	Lys	Asp	Asp	Gln	Gln	Leu	Ser	Pro
2995						3000						3005			
Gly	Phe	Lys	Cys	Lys	Phe	Leu	His	Asp	Ala	Gln	Glu	Tyr	Thr	Leu	Leu
3010						3015						3020			
Leu	Ile	Glu	Thr	Phe	Pro	Glu	Asp	Ser	Ala	Val	Tyr	Thr	Cys	Glu	Ala
3025						3030						3035			3040
Lys	Asn	Asp	Tyr	Gly	Val	Ala	Thr	Thr	Ser	Ala	Ser	Leu	Ser	Val	Glu
			3045						3050						3055
Ile	Pro	Glu	Val	Val	Ser	Pro	Glu	Leu	Glu	Val	Pro	Val	Tyr	Pro	Pro
			3060						3065						3070
Ala	Val	Ile	Val	Pro	Leu	Arg	Asp	Ala	Val	Thr	Ser	Glu	Gly	Gln	Ser
3075						3080						3085			
Ala	Arg	Phe	Gln	Cys	Arg	Val	Thr	Gly	Thr	Asp	Leu	Lys	Val	Ser	Trp
3090						3095						3100			
Tyr	Ser	Lys	Asp	Arg	Glu	Ile	Lys	Pro	Ser	Arg	Phe	Phe	Arg	Met	Thr
3105						3110						3115			3120
Gln	Phe	Glu	Asp	Thr	Tyr	Gln	Leu	Glu	Ile	Ala	Glu	Ala	Tyr	Pro	Glu
			3125						3130						3135
Asp	Glu	Gly	Thr	Tyr	Thr	Phe	Val	Ala	Ser	Asn	Ser	Val	Gly	Gln	Val
			3140						3145						3150
Thr	Ser	Thr	Ala	Ile	Leu	Lys	Leu	Glu	Ala	Pro	Glu	Lys	Ile	Met	Tyr
			3155						3160						3165
Glu	Lys	Leu	Glu	Glu	Glu	Ile	Glu	Met	Glu	Val	Lys	Val	Ala	Pro	Ile
3170						3175						3180			
Leu	Arg	Arg	Arg	Leu	Glu	Pro	Leu	Glu	Val	Ala	Val	Asn	His	Val	Ala
3185						3190						3195			3200
Lys	Phe	Thr	Cys	Glu	Val	Glu	Thr	Thr	Pro	Asn	Val	Lys	Phe	Gln	Trp
			3205						3210						3215
Tyr	Lys	Ala	Gly	Arg	Glu	Ile	Tyr	Asp	Gly	Asp	Lys	Tyr	Ser	Ile	Arg
			3220						3225						3230
Ser	Ser	Asn	Tyr	Leu	Ser	Thr	Leu	Glu	Ile	Pro	Arg	Pro	Gln	Val	Val
3235						3240						3245			
Asp	Cys	Gly	Glu	Tyr	Ser	Cys	Lys	Ala	Ser	Asn	Gln	His	Gly	Ser	Val
3250						3255						3260			
Ser	Ser	Thr	Ala	Phe	Leu	Thr	Val	Thr	Glu	Pro	Pro	Arg	Phe	Ile	Lys
3265						3270						3275			3280
Lys	Leu	Asp	Ser	Ser	Arg	Leu	Val	Lys	Gln	His	Asp	Ser	Thr	Arg	Tyr

3285	3290	3295
Glu Cys Lys Val Gly Gly Ser Pro Glu Ile Lys Val Thr Trp Tyr Lys		
3300	3305	3310
Gly Glu Thr Glu Ile His Pro Ser Glu Lys Tyr Ser Met Ser Phe Val		
3315	3320	3325
Asp Ser Val Ala Val Leu Glu Met His Asn Leu Ser Val Glu Asp Ser		
3330	3335	3340
Gly Asp Tyr Ser Cys Glu Ala Gln Asn Pro Ala Gly Ser Ala Ser Thr		
3345	3350	3355
Ser Thr Ser Leu Lys Val Lys Ala Pro Pro Ala Phe Thr Lys Lys Pro		
3365	3370	3375
His Pro Val Gln Thr Leu Lys Gly Ser Asp Val His Leu Glu Cys Glu		
3380	3385	3390
Leu Gln Gly Thr Pro Pro Phe Gln Ile Ser Trp Tyr Lys Asp Lys Arg		
3395	3400	3405
Glu Ile Arg Ser Ser Lys Lys Tyr Lys Val Met Ser Glu Asn Tyr Leu		
3410	3415	3420
Ala Ser Ile His Ile Leu Asn Val Asp Thr Ala Asp Val Gly Glu Tyr		
3425	3430	3435
His Cys Lys Ala Val Asn Asp Val Gly Ser Asp Ser Cys Ile Gly Ser		
3445	3450	3455
Val Thr Leu Arg Ala Pro Pro Thr Phe Val Lys Lys Leu Ser Asp Val		
3460	3465	3470
Thr Val Val Val Gly Glu Thr Ile Glu Leu Gln Ala Ala Val Glu Gly		
3475	3480	3485
Ala Gln Pro Ile Ser Val Leu Trp Leu Lys Asp Lys Gly Glu Ile Ile		
3490	3495	3500
Arg Glu Ser Glu Asn Leu Trp Ile Ser Tyr Ser Glu Asn Val Ala Ser		
3505	3510	3515
Leu Lys Ile Gly Asn Ala Glu Pro Thr Asn Ala Gly Lys Tyr Ile Cys		
3525	3530	3535
Gln Ile Lys Asn Asp Ala Gly Phe Gln Glu Cys Phe Ala Lys Leu Thr		
3540	3545	3550
Val Leu Glu Pro Ala Val Ile Val Glu Lys Pro Gly Pro Val Lys Val		
3555	3560	3565
Thr Ala Gly Asp Ser Cys Thr Leu Glu Cys Thr Val Asp Gly Thr Pro		
3570	3575	3580
Glu Leu Thr Ala Arg Trp Phe Lys Asp Gly Asn Glu Leu Ser Thr Asp		





[illegible]

<221> VARIANT

<223> Wherein Xaa is any amino acid as defined in the specification.

## <221> VARIANT

<223> Wherein Xaa is any amino acid as defined in the specification.

## <221> VARIANT

<223> Wherein Xaa is any amino acid as defined in the specification.

Met Ile Ser Trp Glu Val Val His Thr Val Phe Leu Phe Ala Leu Leu  
1 5 10 15

Tyr Ser Ser Leu Ala Gln Asp Ala Ser Pro Gln Ser Glu Ile Arg Ala  
20 25 30

Glu Glu Phe Pro Glu Gly Ala Ser Thr Leu Ala Phe Val Phe Asp Val  
35 40 45

Thr Gly Ser Met Tyr Asp Asp Leu Val Gln Val Ile Glu Gly Ala Ser  
50 55 60

Lys Ile Leu Glu Thr Ser Leu Lys Arg Pro Lys Arg Pro Leu Phe Asn  
65 70 75 80

Phe Ala Leu Val Pro Phe His Asp Pro Glu Ile Gly Pro Val Thr Ile  
85 90 95

Thr Thr Asp Pro Lys Lys Phe Gln Tyr Glu Leu Arg Glu Leu Tyr Val  
100 105 110

Gln Gly Gly Gly Asp Cys Pro Glu Met Ser Ile Gly Ala Ile Lys Ile  
115 120 125

Ala Leu Glu Ile Ser Leu Pro Gly Ser Phe Ile Tyr Val Phe Thr Asp  
130 135 140

Ala Arg Ser Lys Asp Tyr Arg Leu Thr His Glu Val Leu Gln Leu Ile  
145 150 155 160

Gln Gln Lys Gln Ser Gln Val Val Phe Val Leu Thr Gly Asp Cys Asp  
165 170 175

Asp Arg Thr His Ile Gly Tyr Lys Val Tyr Glu Glu Ile Ala Ser Thr  
180 185 190

Ser Ser Gly Gln Val Phe His Leu Asp Lys Lys Gln Val Asn Glu Val  
 195 200 205  
 Leu Lys Trp Val Glu Glu Ala Val Gln Ala Ser Lys Val His Leu Leu  
 210 215 220  
 Ser Thr Asp His Leu Glu Gln Ala Val Asn Thr Trp Arg Ile Pro Phe  
 225 230 235 240  
 Asp Pro Ser Leu Lys Glu Val Thr Val Ser Leu Ser Gly Pro Ser Pro  
 245 250 255  
 Met Ile Glu Ile Arg Asn Pro Leu Gly Lys Leu Ile Lys Lys Gly Phe  
 260 265 270  
 Gly Leu His Glu Leu Leu Asn Ile His Asn Ser Ala Lys Val Val Asn  
 275 280 285  
 Val Lys Glu Pro Glu Ala Gly Met Trp Thr Val Lys Thr Ser Ser Ser  
 290 295 300  
 Gly Arg His Ser Val Arg Ile Thr Gly Leu Ser Thr Ile Asp Phe Arg  
 305 310 315 320  
 Ala Gly Phe Ser Arg Lys Pro Thr Leu Asp Phe Lys Lys Thr Val Ser  
 325 330 335  
 Arg Pro Val Gln Gly Ile Pro Thr Tyr Val Leu Leu Asn Thr Ser Gly  
 340 345 350  
 Ile Ser Thr Pro Ala Arg Ile Asp Leu Leu Glu Leu Leu Ser Ile Ser  
 355 360 365  
 Gly Ser Ser Leu Lys Thr Ile Pro Val Lys Tyr Tyr Pro His Arg Lys  
 370 375 380  
 Pro Tyr Gly Ile Trp Asn Ile Ser Asp Phe Val Pro Pro Asn Glu Ala  
 385 390 395 400  
 Phe Phe Leu Lys Val Thr Gly Tyr Asp Lys Asp Asp Tyr Leu Phe Gln  
 405 410 415  
 Arg Val Ser Ser Val Ser Phe Ser Ser Ile Val Pro Asp Ala Pro Lys  
 420 425 430  
 Val Thr Met Pro Glu Lys Thr Pro Gly Tyr Tyr Leu Gln Pro Gly Gln  
 435 440 445  
 Ile Pro Cys Ser Val Asp Ser Leu Leu Pro Phe Thr Leu Ser Phe Val  
 450 455 460  
 Arg Asn Gly Val Thr Leu Gly Val Asp Gln Tyr Leu Lys Glu Ser Ala  
 465 470 475 480  
 Ser Val Ser Leu Asp Ile Ala Lys Val Thr Leu Ser Asp Glu Gly Phe  
 485 490 495

Tyr Glu Cys Ile Ala Val Ser Ser Ala Gly Thr Gly Arg Ala Gln Thr  
 500 505 510  
 Phe Phe Asp Val Ser Glu Pro Pro Pro Val Ile Gln Val Pro Asn Asn  
 515 520 525  
 Val Thr Val Thr Pro Gly Glu Arg Ala Val Leu Thr Cys Leu Ile Ile  
 530 535 540  
 Ser Ala Val Asp Tyr Asn Leu Thr Trp Gln Arg Asn Asp Arg Asp Val  
 545 550 555 560  
 Arg Leu Ala Glu Pro Ala Arg Ile Arg Thr Leu Ala Asn Leu Ser Leu  
 565 570 575  
 Glu Leu Lys Ser Val Lys Phe Asn Asp Ala Gly Glu Tyr His Cys Met  
 580 585 590  
 Val Ser Ser Glu Gly Gly Ser Ser Ala Ala Ser Val Phe Leu Thr Val  
 595 600 605  
 Gln Glu Pro Pro Lys Val Thr Val Met Pro Lys Asn Gln Ser Phe Thr  
 610 615 620  
 Gly Gly Ser Glu Val Ser Ile Met Cys Ser Ala Thr Gly Tyr Pro Lys  
 625 630 635 640  
 Pro Lys Ile Ala Trp Thr Val Asn Asp Met Phe Ile Val Gly Ser His  
 645 650 655  
 Arg Tyr Arg Met Thr Ser Asp Gly Thr Leu Phe Ile Lys Asn Ala Ala  
 660 665 670  
 Pro Lys Asp Ala Gly Ile Tyr Gly Cys Leu Ala Ser Asn Ser Ala Gly  
 675 680 685  
 Thr Asp Lys Gln Asn Ser Thr Leu Arg Tyr Ile Glu Ala Pro Lys Leu  
 690 695 700  
 Met Val Val Gln Ser Glu Leu Leu Val Ala Leu Gly Asp Ile Thr Val  
 705 710 715 720  
 Met Glu Cys Lys Thr Ser Gly Ile Pro Pro Pro Gln Val Lys Trp Phe  
 725 730 735  
 Lys Gly Asp Leu Glu Leu Arg Pro Ser Thr Phe Leu Ile Ile Asp Pro  
 740 745 750  
 Leu Leu Gly Leu Leu Lys Ile Gln Glu Thr Gln Asp Leu Asp Ala Gly  
 755 760 765  
 Asp Tyr Thr Cys Val Ala Ile Asn Glu Ala Gly Arg Ala Thr Gly Lys  
 770 775 780  
 Ile Thr Leu Asp Val Gly Ser Pro Pro Val Phe Ile Gln Glu Pro Ala  
 785 790 795 800



Asp Val Ser Met Glu Ile Gly Ser Asn Val Thr Leu Pro Cys Tyr Val  
 805 810 815  
 Gln Gly Tyr Pro Glu Pro Thr Ile Lys Trp Arg Arg Leu Asp Asn Met  
 820 825 830  
 Pro Ile Phe Ser Arg Pro Phe Ser Val Ser Ser Ile Ser Gln Leu Arg  
 835 840 845  
 Thr Gly Ala Leu Phe Ile Leu Asn Leu Trp Ala Ser Asp Lys Gly Thr  
 850 855 860  
 Tyr Ile Cys Glu Ala Glu Asn Gln Phe Gly Lys Ile Gln Ser Glu Thr  
 865 870 875 880  
 Thr Val Thr Val Thr Gly Leu Val Ala Pro Leu Ile Gly Ile Ser Pro  
 885 890 895  
 Ser Val Ala Asn Val Ile Glu Gly Gln Gln Leu Thr Leu Pro Cys Thr  
 900 905 910  
 Leu Leu Ala Gly Asn Pro Ile Pro Glu Arg Arg Trp Ile Lys Asn Ser  
 915 920 925  
 Ala Met Leu Leu Gln Asn Pro Tyr Ile Thr Val Arg Ser Asp Gly Ser  
 930 935 940  
 Leu His Ile Glu Arg Val Gln Leu Gln Asp Gly Gly Glu Tyr Thr Cys  
 945 950 955 960  
 Val Ala Ser Asn Val Ala Gly Thr Asn Asn Lys Thr Thr Ser Val Val  
 965 970 975  
 Val His Val Leu Pro Thr Ile Gln His Gly Gln Gln Ile Leu Ser Thr  
 980 985 990  
 Ile Glu Gly Ile Pro Val Thr Leu Pro Cys Lys Ala Ser Gly Asn Pro  
 995 1000 1005  
 Lys Pro Ser Val Ile Trp Ser Lys Lys Gly Glu Leu Ile Ser Thr Ser  
 1010 1015 1020  
 Ser Ala Lys Phe Ser Ala Gly Ala Asp Gly Ser Leu Tyr Val Val Ser  
 1025 1030 1035 1040  
 Pro Glu Gly Glu Glu Ser Gly Glu Tyr Val Cys Thr Ala Thr Asn Thr  
 1045 1050 1055  
 Ala Gly Tyr Ala Lys Arg Lys Val Gln Leu Thr Val Tyr Val Arg Pro  
 1060 1065 1070  
 Arg Val Phe Gly Asp Leu Arg Gly Leu Ser Gln Asp Lys Pro Val Glu  
 1075 1080 1085  
 Ile Ser Val Leu Ala Gly Glu Glu Val Thr Leu Pro Cys Glu Val Lys  
 1090 1095 1100

Ser Leu Pro Pro Pro Ile Ile Thr Trp Ala Lys Glu Thr Gln Leu Ile  
 1105 1110 1115 1120  
 Ser Pro Phe Ser Pro Arg His Thr Phe Leu Pro Ser Gly Ser Met Lys  
 1125 1130 1135  
 Ile Thr Glu Thr Arg Thr Ser Asp Ser Gly Met Tyr Leu Cys Val Ala  
 1140 1145 1150  
 Thr Asn Ile Ala Gly Asn Val Thr Gln Ala Val Lys Leu Asn Val His  
 1155 1160 1165  
 Val Pro Pro Lys Ile Gln Arg Gly Pro Lys His Leu Lys Val Gln Val  
 1170 1175 1180  
 Gly Gln Arg Val Asp Ile Pro Cys Asn Ala Gln Gly Thr Pro Leu Pro  
 1185 1190 1195 1200  
 Val Ile Thr Trp Ser Lys Gly Gly Ser Thr Met Leu Val Asp Gly Glu  
 1205 1210 1215  
 His His Val Ser Asn Pro Asp Gly Thr Leu Ser Ile Asp Gln Ala Thr  
 1220 1225 1230  
 Pro Ser Asp Ala Gly Ile Tyr Thr Cys Val Ala Thr Asn Ile Ala Gly  
 1235 1240 1245  
 Thr Asp Glu Thr Glu Ile Thr Leu His Val Gln Glu Pro Pro Thr Val  
 1250 1255 1260  
 Glu Asp Leu Glu Pro Pro Tyr Asn Thr Thr Phe Gln Glu Arg Val Ala  
 1265 1270 1275 1280  
 Asn Gln Arg Ile Glu Phe Pro Cys Pro Ala Lys Gly Thr Pro Lys Pro  
 1285 1290 1295  
 Thr Ile Lys Trp Leu His Asn Gly Arg Glu Leu Thr Gly Arg Glu Pro  
 1300 1305 1310  
 Gly Ile Ser Ile Leu Glu Glu Gly Thr Leu Leu Val Ile Ala Ser Val  
 1315 1320 1325  
 Thr Pro Tyr Asp Asn Gly Glu Tyr Ile Cys Val Ala Val Asn Glu Ala  
 1330 1335 1340  
 Gly Thr Thr Glu Arg Lys Tyr Asn Leu Lys Val His Val Pro Pro Val  
 1345 1350 1355 1360  
 Ile Lys Asp Lys Glu Gln Val Ser Asn Val Ser Val Leu Leu Asn Gln  
 1365 1370 1375  
 Leu Thr Asn Leu Phe Cys Glu Val Glu Gly Thr Pro Ser Pro Ile Ile  
 1380 1385 1390  
 Met Trp Tyr Lys Asp Asn Val Gln Val Thr Glu Ser Ser Thr Ile Gln  
 1395 1400 1405



Val Ala Thr Ser Val Ala Gly Glu Lys Glu Ile Lys Tyr Glu Val Asp  
1715 1720 1725

Val Leu Val Pro Pro Ala Ile Glu Gly Gly Asp Glu Thr Ser Tyr Phe  
1730 1735 1740

Ile Val Met Val Asn Asn Leu Leu Glu Leu Asp Cys His Val Thr Gly  
1745 1750 1755 1760

Ser Pro Pro Pro Thr Ile Met Trp Leu Lys Asp Gly Gln Leu Ile Asp  
1765 1770 1775

Glu Arg Asp Gly Phe Lys Ile Leu Leu Asn Gly Arg Lys Leu Val Ile  
1780 1785 1790

Ala Gln Ala Gln Val Ser Asn Thr Gly Leu Tyr Arg Cys Met Ala Ala  
1795 1800 1805

Asn Thr Ala Gly Asp His Lys Lys Glu Phe Glu Val Thr Val His Val  
1810 1815 1820

Pro Pro Thr Ile Lys Ser Ser Gly Leu Ser Glu Arg Val Val Val Lys  
1825 1830 1835 1840

Tyr Lys Pro Val Ala Leu Gln Cys Ile Ala Asn Gly Ile Pro Asn Pro  
1845 1850 1855

Ser Ile Thr Trp Leu Lys Asp Asp Gln Pro Val Asn Thr Ala Gln Gly  
1860 1865 1870

Asn Leu Lys Ile Gln Ser Ser Gly Arg Val Leu Gln Ile Ala Lys Thr  
1875 1880 1885

Leu Leu Glu Asp Ala Gly Arg Tyr Thr Cys Val Ala Thr Asn Ala Ala  
1890 1895 1900

Gly Glu Thr Gln Gln His Ile Gln Leu His Val His Glu Pro Pro Ser  
1905 1910 1915 1920

Leu Glu Asp Ala Gly Lys Met Leu Asn Glu Thr Val Leu Val Ser Asn  
1925 1930 1935

Pro Val Gln Leu Glu Cys Lys Ala Ala Gly Asn Pro Val Pro Val Ile  
1940 1945 1950

Thr Trp Tyr Lys Asp Asn Cys Leu Ser Gly Ser Thr Ser Met Thr  
1955 1960 1965

Phe Leu Asn Arg Gly Gln Ile Ile Asp Ile Glu Ser Ala Gln Ile Ser  
1970 1975 1980

Asp Ala Gly Ile Tyr Lys Cys Val Ala Ile Asn Ser Ala Gly Ala Thr  
1985 1990 1995 2000

Glu Leu Phe Tyr Ser Leu Gln Val His Val Ala Pro Ser Ile Ser Gly  
2005 2010 2015



Pro Pro Pro Thr Val Thr Trp Met Lys Asp Gly His Pro Leu Ile Lys  
2325 2330 2335

Ala Lys Gly Val Glu Ile Leu Asp Glu Gly His Ile Leu Gln Leu Lys  
2340 2345 2350

Asn Ile His Val Ser Asp Thr Gly Arg Tyr Val Cys Val Ala Val Asn  
2355 2360 2365

Val Ala Gly Met Thr Asp Lys Lys Tyr Asp Leu Ser Val His Ala Pro  
2370 2375 2380

Pro Ser Ile Ile Gly Asn His Arg Ser Pro Glu Asn Ile Ser Val Val  
2385 2390 2395 2400

Glu Lys Asn Ser Val Ser Leu Thr Cys Glu Ala Ser Gly Ile Pro Leu  
2405 2410 2415

Pro Ser Thr Thr Trp Phe Lys Asp Gly Trp Pro Val Ser Leu Ser Asn  
2420 2425 2430

Ser Val Arg Ile Leu Ser Gly Gly Arg Met Leu Arg Leu Met Gln Thr  
2435 2440 2445

Thr Met Glu Asp Ala Gly Gln Tyr Thr Cys Val Val Arg Asn Ala Ala  
2450 2455 2460

Gly Glu Glu Arg Lys Ile Phe Gly Leu Ser Val Leu Val Pro Pro His  
2465 2470 2475 2480

Ile Val Gly Glu Asn Thr Leu Glu Asp Val Lys Val Lys Glu Lys Gln  
2485 2490 2495

Ser Val Thr Leu Thr Cys Glu Val Thr Gly Asn Pro Val Pro Glu Ile  
2500 2505 2510

Thr Trp His Lys Asp Gly Gln Pro Leu Gln Glu Asp Glu Ala His His  
2515 2520 2525

Ile Ile Ser Gly Gly Arg Phe Leu Gln Ile Thr Asn Val Gln Val Pro  
2530 2535 2540

His Thr Gly Arg Tyr Thr Cys Leu Ala Ser Ser Pro Ala Gly His Lys  
2545 2550 2555 2560

Ser Arg Ser Phe Ser Leu Asn Val Phe Val Ser Pro Thr Ile Ala Gly  
2565 2570 2575

Val Gly Ser Asp Gly Asn Pro Glu Asp Val Thr Val Ile Leu Asn Ser  
2580 2585 2590

Pro Thr Ser Leu Val Cys Glu Ala Tyr Ser Tyr Pro Pro Ala Thr Ile  
2595 2600 2605

Thr Trp Phe Lys Asp Gly Thr Pro Leu Glu Ser Asn Arg Asn Ile Arg  
2610 2615 2620



Leu Gln Ile Leu Asn Thr Gln Ile Thr Asp Ile Gly Arg Tyr Val Cys  
 2930 2935 2940

Val Ala Glu Asn Thr Ala Gly Ser Ala Lys Lys Tyr Phe Asn Leu Asn  
 2945 2950 2955 2960

Val His Val Pro Pro Ser Val Ile Gly Pro Lys Ser Glu Asn Leu Thr  
 2965 2970 2975

Val Val Val Asn Asn Phe Ile Ser Leu Thr Cys Glu Val Ser Gly Phe  
 2980 2985 2990

Pro Pro Pro Asp Leu Ser Trp Leu Lys Asn Xaa Gln Pro Ile Lys Leu  
 2995 3000 3005

Asn Thr Asn Thr Leu Ile Val Pro Gly Gly Arg Thr Leu Gln Ile Ile  
 3010 3015 3020

Arg Ala Lys Val Ser Asp Gly Gly Glu Tyr Thr Cys Ile Ala Ile Asn  
 3025 3030 3035 3040

Xaa Ala Gly Glu Ser Lys Lys Lys Phe Ser Leu Thr Val Tyr Val Pro  
 3045 3050 3055

Pro Ser Ile Lys Asp His Asp Ser Glu Ser Leu Ser Val Val Asn Val  
 3060 3065 3070

Arg Glu Gly Thr Ser Val Ser Leu Glu Cys Glu Ser Asn Ala Val Pro  
 3075 3080 3085

Pro Pro Val Ile Thr Trp Tyr Lys Asn Gly Arg Met Ile Thr Glu Ser  
 3090 3095 3100

Thr His Val Glu Ile Leu Ala Asp Gly Gln Met Leu His Ile Lys Lys  
 3105 3110 3115 3120

Ala Glu Val Ser Asp Thr Gly Gln Tyr Val Cys Arg Ala Ile Asn Val  
 3125 3130 3135

Ala Gly Arg Asp Asp Lys Asn Phe His Leu Asn Val Tyr Val Pro Pro  
 3140 3145 3150

Ser Ile Glu Gly Pro Glu Arg Glu Val Ile Val Glu Thr Ile Ser Asn  
 3155 3160 3165

Pro Val Thr Leu Thr Cys Asp Ala Thr Gly Ile Pro Pro Pro Thr Ile  
 3170 3175 3180

Ala Trp Leu Lys Asn Tyr Lys Arg Ile Glu Asn Ser Asp Ser Leu Glu  
 3185 3190 3195 3200

Val Arg Ile Leu Ser Gly Gly Ser Lys Leu Gln Ile Ala Arg Ser Gln  
 3205 3210 3215

His Ser Asp Ser Gly Asn Tyr Thr Cys Ile Ala Ser Asn Met Glu Gly  
 3220 3225 3230



Lys Ala Gln Lys Tyr Tyr Phe Leu Ser Ile Gln Val Pro Pro Ser Val  
3235 3240 3245

Ala Gly Ala Glu Ile Pro Ser Asp Val Ser Val Leu Leu Gly Glu Asn  
3250 3255 3260

Val Glu Leu Val Cys Asn Ala Asn Gly Ile Pro Thr Pro Leu Ile Gln  
3265 3270 3275 3280

Trp Leu Lys Asp Gly Lys Pro Ile Ala Ser Gly Glu Thr Glu Arg Ile  
3285 3290 3295

Arg Val Ser Ala Asn Gly Ser Thr Leu Asn Ile Tyr Gly Ala Leu Thr  
3300 3305 3310

Ser Asp Thr Gly Lys Tyr Thr Cys Val Ala Thr Asn Pro Ala Gly Glu  
3315 3320 3325

Glu Asp Arg Ile Phe Asn Leu Asn Val Tyr Val Thr Pro Thr Ile Arg  
3330 3335 3340

Gly Asn Lys Asp Glu Ala Glu Lys Leu Met Thr Tyr Val Asp Thr Ser  
3345 3350 3355 3360

Ile Asn Ile Glu Cys Arg Xaa Thr Gly Thr Pro Pro Pro Gln Ile Asn  
3365 3370 3375

Trp Leu Lys Asn Gly Leu Pro Leu Pro Leu Ser Ser His Ile Arg Leu  
3380 3385 3390

Leu Ala Ala Gly Gln Val Ile Arg Ile Val Arg Ala Gln Val Ser Asp  
3395 3400 3405

Val Ala Val Tyr Thr Cys Val Ala Ser Asn Arg Ala Gly Val Asp Asn  
3410 3415 3420

Lys His Tyr Asn Leu Gln Val Phe Ala Pro Pro Asn Met Asp Asn Ser  
3425 3430 3435 3440

Met Gly Thr Glu Glu Ile Thr Val Leu Lys Gly Ser Ser Thr Ser Met  
3445 3450 3455

Ala Cys Ile Thr Asp Gly Thr Pro Ala Pro Ser Met Ala Trp Leu Arg  
3460 3465 3470

Asp Gly Gln Pro Leu Gly Leu Asp Ala His Leu Thr Val Ser Thr His  
3475 3480 3485

Gly Met Val Leu Gln Leu Leu Lys Ala Glu Thr Glu Asp Ser Gly Lys  
3490 3495 3500

Tyr Thr Cys Ile Ala Ser Asn Glu Ala Gly Glu Val Ser Lys His Phe  
3505 3510 3515 3520

Ile Leu Lys Val Leu Glu Pro Pro His Ile Asn Gly Ser Glu Glu His  
3525 3530 3535





Gly His Tyr Thr Cys Met Ala Ala Asn Val Ala Gly Ser Ser Ser Thr  
 4145 4150 4155 4160  
 Ser Thr Lys Leu Thr Val His Val Pro Pro Arg Ile Arg Ser Thr Glu  
 4165 4170 4175  
 Gly His Tyr Thr Val Asn Glu Asn Ser Gln Ala Ile Leu Pro Cys Val  
 4180 4185 4190  
 Ala Asp Gly Ile Pro Thr Pro Ala Ile Asn Trp Lys Lys Asp Asn Val  
 4195 4200 4205  
 Leu Leu Ala Asn Leu Leu Gly Lys Tyr Thr Ala Glu Pro Tyr Gly Glu  
 4210 4215 4220  
 Leu Ile Leu Glu Asn Val Val Leu Glu Asp Ser Gly Phe Tyr Thr Cys  
 4225 4230 4235 4240  
 Val Ala Asn Asn Ala Ala Gly Glu Asp Thr His Thr Val Ser Leu Thr  
 4245 4250 4255  
 Val His Val Leu Pro Thr Phe Thr Glu Leu Pro Gly Asp Val Ser Leu  
 4260 4265 4270  
 Asn Lys Gly Glu Gln Leu Arg Leu Ser Cys Lys Ala Thr Gly Ile Pro  
 4275 4280 4285  
 Leu Pro Lys Leu Thr Trp Thr Phe Asn Asn Asn Ile Ile Pro Ala His  
 4290 4295 4300  
 Phe Asp Ser Val Asn Gly His Ser Glu Leu Val Ile Glu Arg Val Ser  
 4305 4310 4315 4320  
 Lys Glu Asp Ser Gly Thr Tyr Val Cys Thr Ala Glu Asn Ser Val Gly  
 4325 4330 4335  
 Phe Val Lys Ala Thr Gly Phe Val Tyr Val Lys Glu Pro Pro Val Phe  
 4340 4345 4350  
 Lys Gly Asp Tyr Pro Ser Asn Trp Ile Glu Pro Leu Gly Gly Asn Ala  
 4355 4360 4365  
 Ile Leu Asn Cys Glu Val Lys Gly Asp Pro Thr Pro Thr Ile Gln Trp  
 4370 4375 4380  
 Asn Arg Lys Gly Val Asp Ile Glu Ile Ser His Arg Ile Arg Gln Leu  
 4385 4390 4395 4400  
 Gly Asn Gly Ser Leu Ala Ile Tyr Gly Thr Val Asn Glu Asp Ala Gly  
 4405 4410 4415  
 Asp Tyr Thr Cys Val Ala Thr Asn Glu Ala Gly Val Val Glu Arg Ser  
 4420 4425 4430  
 Met Ser Leu Thr Leu Gln Ser Pro Pro Ile Ile Thr Leu Glu Pro Val  
 4435 4440 4445



Asp Pro Cys Pro Thr His Gly Asn Trp Ser Pro Trp Ser Gly Trp Gly  
4755 4760 4765

Thr Cys Ser Arg Thr Cys Asn Gly Gly Gln Met Arg Arg Tyr Arg Thr  
4770 4775 4780

Cys Asp Asn Pro Pro Pro Ser Asn Gly Gly Arg Ala Cys Gly Gly Pro  
4785 4790 4795 4800

Asp Ser Gln Ile Gln Arg Cys Asn Thr Asp Met Cys Pro Val Asp Gly  
4805 4810 4815

Ser Trp Gly Ser Trp His Ser Trp Ser Gln Cys Ser Ala Ser Cys Gly  
4820 4825 4830

Gly Gly Glu Lys Thr Arg Lys Arg Leu Cys Asp His Pro Val Pro Val  
4835 4840 4845

Lys Gly Gly Arg Pro Cys Pro Gly Asp Thr Thr Gln Val Thr Arg Cys  
4850 4855 4860

Asn Val Gln Ala Cys Pro Gly Gly Pro Gln Arg Ala Arg Gly Ser Val  
4865 4870 4875 4880

Ile Gly Asn Ile Asn Asp Val Glu Phe Gly Ile Ala Phe Leu Asn Ala  
4885 4890 4895

Thr Ile Thr Asp Ser Pro Asn Ser Asp Thr Arg Ile Ile Arg Ala Lys  
4900 4905 4910

Ile Thr Asn Val Pro Arg Ser Leu Gly Ser Ala Met Arg Lys Ile Val  
4915 4920 4925

Ser Ile Leu Asn Pro Ile Tyr Trp Thr Thr Ala Lys Glu Ile Gly Glu  
4930 4935 4940

Ala Val Asn Gly Phe Thr Leu Thr Asn Ala Val Phe Lys Arg Glu Thr  
4945 4950 4955 4960

Gln Val Glu Phe Ala Thr Gly Glu Ile Leu Gln Met Ser His Ile Ala  
4965 4970 4975

Arg Gly Leu Asp Ser Asp Gly Ser Leu Leu Leu Asp Ile Val Val Ser  
4980 4985 4990

Gly Tyr Val Leu Gln Leu Gln Ser Pro Ala Glu Val Thr Val Lys Asp  
4995 5000 5005

Tyr Thr Glu Asp Tyr Ile Gln Thr Gly Pro Gly Gln Leu Tyr Ala Tyr  
5010 5015 5020

Ser Thr Arg Leu Phe Thr Ile Asp Gly Ile Ser Ile Pro Tyr Thr Trp  
5025 5030 5035 5040

Asn His Thr Val Phe Tyr Asp Gln Ala Gln Gly Arg Met Pro Phe Leu  
5045 5050 5055

Val Glu Thr Leu His Ala Ser Ser Val Glu Ser Asp Tyr Asn Gln Ile  
5060 5065 5070

Glu Glu Thr Leu Gly Phe Lys Ile His Ala Ser Ile Ser Lys Gly Asp  
5075 5080 5085

Arg Ser Asn Gln Cys Pro Ser Gly Phe Thr Leu Asp Ser Val Gly Pro  
5090 5095 5100

Phe Cys Ala Asp Glu Asp Glu Cys Ala Ala Gly Asn Pro Cys Ser His  
5105 5110 5115 5120

Ser Cys His Asn Ala Met Gly Thr Tyr Tyr Cys Ser Cys Pro Lys Gly  
5125 5130 5135

Leu Thr Ile Ala Ala Asp Gly Arg Thr Cys Gln Asp Ile Asp Glu Cys  
5140 5145 5150

Ala Leu Gly Arg His Thr Cys His Ala Gly Gln Asp Cys Asp Asn Thr  
5155 5160 5165

Ile Gly Ser Tyr Arg Cys Val Val Arg Cys Gly Ser Gly Phe Arg Arg  
5170 5175 5180

Thr Ser Asp Gly Leu Ser Cys Gln Asp Ile Asn Glu Cys Gln Glu Ser  
5185 5190 5195 5200

Ser Pro Cys His Gln Arg Cys Phe Asn Ala Ile Gly Ser Phe His Cys  
5205 5210 5215

Gly Cys Glu Pro Gly Tyr Gln Leu Lys Gly Arg Lys Cys Met Asp Val  
5220 5225 5230

Asn Glu Cys Arg Gln Asn Val Cys Arg Pro Asp Gln His Cys Lys Asn  
5235 5240 5245

Thr Arg Gly Gly Tyr Lys Cys Ile Asp Leu Cys Pro Asn Gly Met Thr  
5250 5255 5260

Lys Ala Glu Asn Gly Thr Cys Ile Asp Ile Asp Glu Cys Lys Asp Gly  
5265 5270 5275 5280

Thr His Gln Cys Arg Tyr Asn Gln Ile Cys Glu Asn Thr Arg Ser Ser  
5285 5290 5295

Tyr Arg Cys Val Cys Pro Arg Gly Tyr Arg Ser Gln Gly Val Gly Arg  
5300 5305 5310

Pro Cys Met Asp Ile Asp Glu Cys Glu Gln Val Pro Lys Pro Cys Ala  
5315 5320 5325

His Gln Cys Ser Asn Thr Pro Gly Ser Phe Lys Cys Ile Cys Pro Pro  
5330 5335 5340

Gly Gln His Leu Leu Gly Asp Gly Lys Ser Cys Ala Gly Leu Glu Arg  
5345 5350 5355 5360

Leu Pro Asn Tyr Gly Thr Gln Tyr Ser Ser Tyr Asn Leu Ala Arg Phe  
5365 5370 5375

Ser Pro Val Arg Asn Asn Tyr Gln Pro Gln Gln His Tyr Arg Gln Tyr  
5380 5385 5390

Ser His Leu Tyr Ser Ser Tyr Ser Glu Tyr Arg Asn Ser Arg Thr Ser  
5395 5400 5405

Leu Ser Arg Thr Arg Arg Thr Ile Arg Lys Thr Cys Pro Glu Gly Ser  
5410 5415 5420

Glu Ala Ser His Asp Thr Cys Val Asp Ile Asp Glu Cys Glu Asn Thr  
5425 5430 5435 5440

Asp Ala Cys Gln His Glu Cys Lys Asn Thr Phe Gly Ser Tyr Gln Cys  
5445 5450 5455

Ile Cys Pro Pro Gly Tyr Gln Leu Thr His Asn Gly Lys Thr Cys Gln  
5460 5465 5470

Asp Ile Asp Glu Cys Leu Glu Gln Asn Val His Cys Gly Pro Asn Arg  
5475 5480 5485

Met Cys Phe Asn Met Arg Gly Ser Tyr Gln Cys Ile Asp Thr Pro Cys  
5490 5495 5500

Pro Pro Asn Tyr Gln Arg Asp Pro Val Ser Gly Phe Cys Leu Lys Asn  
5505 5510 5515 5520

Cys Pro Pro Asn Asp Leu Glu Cys Ala Leu Ser Pro Tyr Ala Leu Glu  
5525 5530 5535

Tyr Lys Leu Val Ser Leu Pro Phe Gly Ile Ala Thr Asn Gln Asp Leu  
5540 5545 5550

Ile Arg Leu Val Ala Tyr Thr Gln Asp Gly Val Met His Pro Arg Thr  
5555 5560 5565

Thr Phe Leu Met Val Asp Glu Glu Gln Thr Val Pro Phe Ala Leu Arg  
5570 5575 5580

Asp Glu Asn Leu Lys Gly Val Val Tyr Thr Thr Arg Pro Leu Arg Glu  
5585 5590 5595 5600

Ala Glu Thr Tyr Arg Met Arg Val Arg Ala Ser Ser Tyr Ser Ala Asn  
5605 5610 5615

Gly Thr Ile Glu Tyr Gln Thr Thr Phe Ile Val Tyr Ile Ala Val Ser  
5620 5625 5630

Ala Tyr Pro Tyr  
5635

<210> 94  
<211> 63



<212> PRT  
<213> Homo sapiens

<400> 94  
Leu Glu Gly Glu Ser Val Thr Leu Thr Cys Pro Ala Ser Gly Asp Pro  
1 5 10 15  
Val Pro Asn Ile Thr Trp Leu Lys Asp Gly Lys Pro Leu Pro Glu Ser  
20 25 30  
Arg Val Val Ala Ser Gly Ser Thr Leu Thr Ile Lys Asn Val Ser Leu  
35 40 45  
Glu Asp Ser Gly Leu Tyr Thr Cys Val Ala Arg Asn Ser Val Gly  
50 55 60

<210> 95  
<211> 81  
<212> PRT  
<213> Homo sapiens

<400> 95  
Val Lys Glu Gly Glu Ser Val Thr Leu Ser Cys Glu Ala Ser Gly Asn  
1 5 10 15  
Pro Pro Pro Thr Val Thr Trp Tyr Lys Gln Gly Gly Lys Leu Leu Ala  
20 25 30  
Glu Ser Gly Arg Phe Ser Val Ser Arg Ser Gly Gly Asn Ser Thr Leu  
35 40 45  
Thr Ile Ser Asn Val Thr Pro Glu Asp Ser Gly Thr Tyr Thr Cys Ala  
50 55 60  
Ala Thr Asn Ser Ser Gly Ser Ala Ser Ser Gly Thr Thr Leu Thr Val  
65 70 75 80  
Leu

<210> 96  
<211> 629  
<212> PRT  
<213> Mus musculus

<400> 96  
Gln Ala Ala Arg Gly Arg Thr Arg Lys Gly Lys Tyr Cys Leu Gln Leu  
1 5 10 15  
Ser Pro Phe Ile Leu Trp Phe Leu Arg Leu Asp Asn Leu Ile Phe His  
20 25 30  
Pro Glu Lys Ala Glu Val Leu Ala Val Leu Asp Trp Glu Leu Ser Thr  
35 40 45

Leu Gly Asp Pro Phe Ala Asp Val Ala Tyr Ser Cys Leu Ala Tyr Tyr  
 50 55 60  
 Leu Pro Ser Ser Phe Pro Ile Leu Arg Gly Phe Arg Asp Gln Asp Val  
 65 70 75 80  
 Thr Lys Leu Gly Ile Pro Thr Val Glu Glu Tyr Phe Arg Met Tyr Cys  
 85 90 95  
 Leu Asn Met Gly Ile Pro Pro Ile Asp Asn Trp Asn Phe Tyr Met Ala  
 100 105 110  
 Phe Ser Phe Phe Arg Val Ala Ala Ile Leu Gln Gly Val Tyr Lys Arg  
 115 120 125  
 Ser Leu Thr Gly Gln Ala Ser Ser Ala Thr Ala Gln Gln Ser Gly Lys  
 130 135 140  
 Leu Thr Glu Ser Met Ala Glu Leu Ala Trp Asp Phe Ala Thr Lys Glu  
 145 150 155 160  
 Gly Phe Arg Val Phe Lys Glu Met Pro Ala Thr Lys Thr Leu Ser Arg  
 165 170 175  
 Ser Tyr His Ala Trp Ala Gly Pro Arg Ser Pro Arg Thr Pro Lys Gly  
 180 185 190  
 Val Arg Gly His Ser Thr Val Ala Ala Ala Ser Pro Ser His Glu Ala  
 195 200 205  
 Lys Gly Gly Leu Val Ile Ser Pro Glu Gly Leu Ser Pro Ala Val Arg  
 210 215 220  
 Lys Leu Tyr Glu Gln Leu Val Gln Phe Ile Glu Gln Lys Val Tyr Pro  
 225 230 235 240  
 Leu Glu Pro Glu Leu Gln Arg His Gln Ala Ser Ala Asp Arg Trp Ser  
 245 250 255  
 Pro Ser Pro Leu Ile Glu Asp Leu Lys Glu Lys Ala Lys Ala Glu Gly  
 260 265 270  
 Leu Trp Asn Leu Phe Leu Pro Leu Glu Thr Asp Pro Glu Lys Lys Tyr  
 275 280 285  
 Gly Ala Gly Leu Thr Asn Val Glu Tyr Ala His Leu Cys Glu Val Met  
 290 295 300  
 Gly Met Ser Leu Tyr Ala Ser Glu Ile Phe Asn Cys Ser Ala Pro Asp  
 305 310 315 320  
 Thr Gly Asn Met Glu Ile Leu Val Arg Tyr Gly Thr Glu Glu Gln Lys  
 325 330 335  
 Ala Arg Trp Leu Val Pro Leu Leu Glu Gly Arg Ile Arg Ser Cys Phe  
 340 345 350

Ala Met Thr Glu Pro Gln Val Ala Ser Ser Asp Ala Ser Asn Ile Glu  
355 360 365

Ala Ser Ile Lys Glu Glu Asp Gly Cys Tyr Val Ile Asn Gly His Lys  
370 375 380

Trp Trp Thr Ser Gly Ile Leu Asp Pro Arg Cys Lys Leu Cys Val Phe  
385 390 395 400

Met Gly Lys Thr Asp Pro Gln Ala Pro Arg His Gln Gln Gln Ser Met  
405 410 415

Leu Leu Val Pro Met Asp Ser Pro Gly Ile Thr Val Ile Arg Pro Leu  
420 425 430

Ser Val Phe Gly Leu Glu Asp Pro Pro Gly Gly His Gly Glu Val Arg  
435 440 445

Phe Lys Asp Val Arg Val Pro Lys Glu Asn Ile Leu Leu Gly Pro Gly  
450 455 460

Arg Gly Phe Glu Ile Ala Gln Gly Arg Leu Gly Pro Gly Arg Ile His  
465 470 475 480

His Cys Met Arg Leu Ile Gly Tyr Ser Glu Arg Ala Leu Ala Leu Met  
485 490 495

Lys Thr Arg Val Met Ser Arg Thr Ala Phe Gly Lys Pro Leu Val Glu  
500 505 510

Gln Gly Thr Ile Leu Ala Asp Ile Ala Arg Ser Arg Val Glu Ile Glu  
515 520 525

Gln Ala Arg Leu Leu Val Leu Lys Ala Ala His Leu Met Asp Val Ala  
530 535 540

Gly Asn Lys Thr Ala Ala Leu Asp Ile Ala Met Ile Lys Met Val Val  
545 550 555 560

Pro Ser Met Ala Tyr His Val Ile Asp Arg Ala Ile Gln Ala Phe Gly  
565 570 575

Ala Ala Gly Leu Ser Ser Asp Tyr Pro Leu Ala Gln Phe Phe Gly Trp  
580 585 590

Ala Arg Ala Leu Arg Phe Ala Asp Gly Pro Asp Glu Val His Gln Leu  
595 600 605

Thr Val Ala Lys Met Glu Leu Lys Asn Gln Ser Arg Met Gln Glu Pro  
610 615 620

Ala Val Pro Arg Val  
625

<210> 97  
<211> 455

<212> PRT

<213> Mus musculus

<400> 97

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Met Glu Gln Arg Val Tyr Pro Ala Glu Pro Glu Leu Gln Ser His Gln
 1          5          10          15

Ala Ser Ala Ala Arg Trp Ser Pro Ser Pro Leu Ile Glu Asp Leu Lys
          20          25          30

Glu Lys Ala Lys Ala Glu Gly Leu Trp Asn Leu Phe Leu Pro Leu Glu
          35          40          45

Ala Asp Pro Glu Lys Lys Tyr Gly Ala Gly Leu Thr Asn Val Glu Tyr
          50          55          60

Ala His Leu Cys Glu Leu Met Gly Thr Ser Leu Tyr Ala Pro Glu Val
          65          70          75          80

Cys Asn Cys Ser Ala Pro Asp Thr Gly Asn Met Glu Leu Leu Val Arg
          85          90          95

Tyr Gly Thr Glu Ala Gln Lys Ala Arg Trp Leu Ile Pro Leu Leu Glu
          100          105          110

Gly Lys Ala Arg Ser Cys Phe Ala Met Thr Glu Pro Gln Val Ala Ser
          115          120          125

Ser Asp Ala Thr Asn Ile Glu Ala Ser Ile Arg Glu Glu Asp Ser Phe
          130          135          140

Tyr Val Ile Asn Gly His Lys Trp Trp Ile Thr Gly Ile Leu Asp Pro
          145          150          155          160

Arg Cys Gln Leu Cys Val Phe Met Gly Lys Thr Asp Pro His Ala Pro
          165          170          175

Arg His Arg Gln Gln Ser Val Leu Leu Val Pro Met Asp Thr Pro Gly
          180          185          190

Ile Lys Ile Ile Arg Pro Leu Thr Val Tyr Gly Leu Glu Asp Ala Pro
          195          200          205

Gly Gly His Gly Glu Val Arg Phe Glu His Val Arg Val Pro Lys Glu
          210          215          220

Asn Met Val Leu Gly Pro Gly Arg Gly Phe Glu Ile Ala Gln Gly Arg
          225          230          235          240

Leu Gly Pro Gly Arg Ile His His Cys Met Arg Leu Ile Gly Phe Ser
          245          250          255

Glu Arg Ala Leu Ala Leu Met Lys Ala Arg Val Lys Ser Arg Leu Ala
          260          265          270

Phe Gly Lys Pro Leu Val Glu Gln Gly Thr Val Leu Ala Asp Ile Ala
          275          280          285

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Gln Ser Arg Val Glu Ile Glu Gln Ala Arg Leu Leu Val Leu Arg Ala  
 290 295 300  
 Ala His Leu Met Asp Leu Ala Gly Asn Lys Ala Ala Ala Leu Asp Ile  
 305 310 315 320  
 Ala Met Ile Lys Met Val Ala Pro Ser Met Ala Ser Arg Val Ile Asp  
 325 330 335  
 Arg Ala Ile Gln Lys Thr Ser Leu Gln Glu Ala Trp Ser Leu Phe Gln  
 340 345 350  
 Ala Arg Arg Arg Gly Phe Ala Glu Gly Gln Gly Gly Ser Gly Thr Glu  
 355 360 365  
 Ser Gly Lys Leu Val Phe Arg Leu Ser Val Pro Gly Trp Ala Gly Thr  
 370 375 380  
 Val Thr Ser Leu Gln Pro Phe Ser Pro Ser Leu Ser Ala Cys Gly Asn  
 385 390 395 400  
 Leu Asp Thr Phe Trp Glu Ala Ser Gln Gly Cys Gly Thr Cys Leu Leu  
 405 410 415  
 Trp Gln Leu Gln Gly Ser Cys Leu Ala Ser Leu Val Ser Arg Gly Ala  
 420 425 430  
 Ala Thr Ala Gly Gly Gly Leu Glu Thr Gln Asp Leu Gly Ala Trp Glu  
 435 440 445  
 Asn Gly Met Gln Pro Thr Leu  
 450 455

<210> 98  
 <211> 415  
 <212> PRT  
 <213> *Deinococcus radiodurans*

<400> 98  
 Met Thr Met Phe Asp Thr Thr Pro Arg Ala Gln Asp Leu Arg Glu Arg  
 1 5 10 15  
 Leu Leu Arg Phe Met Asp Thr Tyr Ile Tyr Pro Asn Glu Ala Glu Phe  
 20 25 30  
 His Arg Gln Val Glu Ser Gly Glu Arg Trp Ala Pro Val Glu Leu Ile  
 35 40 45  
 Glu Glu Leu Lys Pro Lys Ala Arg Ala Glu Gly Leu Trp Asn Leu Phe  
 50 55 60  
 Leu Pro Pro Ala Ser Asp Pro Glu Gly Lys Phe Gly Ala Gly Leu Thr  
 65 70 75 80  
 Asn Leu Glu Tyr Ala Gly Leu Cys Glu Ile Met Gly Arg Val Trp Trp

**SECRET**

Ala	Pro	Glu	Val	Phe	Asn	Cys	Asn	Ala	Pro	Asp	Thr	Gly	Asn	Met	Glu	
				100					105						110	
Val	Ile	Ala	Arg	Tyr	Gly	Thr	Pro	Glu	Gln	Gln	Glu	Gln	Trp	Leu	Leu	
		115					120					125				
Pro	Leu	Leu	Asn	Gly	Glu	Ile	Arg	Ser	Ala	Phe	Ser	Met	Thr	Glu	Pro	
	130					135					140					
Gln	Val	Ala	Ser	Ser	Asp	Ala	Thr	Asn	Ile	Glu	Ala	Gln	Ile	Val	Arg	
145					150					155					160	
Asp	Gly	Asp	Glu	Tyr	Val	Ile	Asn	Gly	Arg	Lys	Trp	Trp	Ser	Ser	Gly	
				165					170					175		
Ala	Gly	Asp	Ser	Arg	Cys	Lys	Val	Ser	Ile	Phe	Met	Gly	Lys	Thr	Asp	
			180					185					190			
Pro	Gln	Ala	Pro	Arg	His	Leu	Gln	Gln	Ser	Met	Ile	Leu	Val	Pro	Phe	
		195					200					205				
Asp	Ala	Pro	Gly	Val	Lys	Ile	Glu	Arg	Ala	Leu	Gln	Val	Phe	Gly	Phe	
	210					215					220					
Asp	Asp	Ala	Pro	His	Gly	His	Met	Glu	Met	Ser	Phe	Asp	Asn	Val	Arg	
225					230					235					240	
Val	Pro	Val	Thr	Asn	Met	Leu	Leu	Gly	Glu	Gly	Arg	Gly	Phe	Glu	Ile	
				245					250					255		
Ala	Gln	Gly	Arg	Leu	Gly	Pro	Gly	Arg	Ile	His	His	Cys	Met	Arg	Leu	
			260					265					270			
Val	Gly	Gln	Ala	Glu	Arg	Ala	Leu	Glu	Leu	Met	Ile	Glu	Arg	Ser	Ala	
		275					280					285				
Gln	Arg	Ile	Ala	Phe	Gly	Lys	Pro	Leu	Ala	Leu	His	Gln	His	Thr	Arg	
	290					295					300					
Glu	Leu	Ile	Ala	Gln	Ser	Arg	Met	Glu	Ile	Asp	Gln	Ala	Arg	Leu	Leu	
305					310					315					320	
Thr	Leu	Lys	Ala	Ala	His	Thr	Met	Asp	Thr	Val	Gly	Asn	Lys	Asp	Ala	
				325					330					335		
Lys	Gly	Glu	Ile	Ala	Ala	Ile	Lys	Val	Val	Ala	Pro	Asn	Met	Ala	Leu	
			340					345					350			
Arg	Val	Ile	Asp	Arg	Ala	Ile	Gln	Val	Tyr	Gly	Gly	Ala	Gly	Val	Cys	
		355					360					365				
Gln	Asp	Thr	Pro	Leu	Ala	Met	Met	Tyr	Ala	Gln	Ala	Arg	Thr	Leu	Arg	
	370					375					380					
Leu	Ala	Asp	Gly	Pro	Asp	Ile	Val	His	Thr	Glu	Thr	Val	Ala	Lys	Glu	



Ile Leu Gly Glu Gly Arg Gly Phe Glu Ile Ala Gln Gly Arg Leu Gly  
245 250 255  
Pro Gly Arg Ile His His Cys Met Arg Ser Ile Gly Met Ala Glu Arg  
260 265 270  
Ala Leu Glu Leu Met Cys Lys Arg Ala Val Ser Arg Thr Ala Phe Gly  
275 280 285  
Lys Pro Leu Ala Arg Leu Gly Gly Asn Ile Asp His Ile Ala Asp Ser  
290 295 300  
Arg Met Glu Ile Asn Met Ala Arg Leu Leu Thr Leu Gln Ala Ala Tyr  
305 310 315 320  
Met Met Asp Thr Val Gly Asn Lys Ile Ala Gln Ser Glu Ile Ala Gln  
325 330 335  
Ile Lys Val Val Ala Pro Asn Val Ala Leu Lys Val Ile Asp Arg Ala  
340 345 350  
Ile Gln Met His Gly Gly Ala Gly Val Ser Asn Asp Phe Pro Leu Ala  
355 360 365  
Tyr Trp Tyr Ala Met Gln Arg Thr Leu Arg Leu Ala Asp Gly Pro Asp  
370 375 380  
Glu Val His Arg Ala Ala Ile Gly Lys Phe Glu Leu Gly Lys Tyr Val  
385 390 395 400  
Pro Arg Glu Met Leu Arg Ser Ser Arg  
405

<210> 100  
<211> 423  
<212> PRT  
<213> Arabidopsis thaliana

<400> 100  
Met Asp Ala Val Gln Arg Asp Val Ser Pro Ser Tyr Glu Ser Leu Val  
1 5 10 15  
Asp Gly Ser Gly Arg Phe Ile Pro Asn Arg Lys Val Leu Glu Leu Arg  
20 25 30  
Gln Lys Leu Ile Lys Phe Met Glu Thr His Ile Tyr Pro Met Glu Asn  
35 40 45  
Glu Phe Ser Lys Leu Ala Gln Ser Asp Met Arg Trp Thr Val His Pro  
50 55 60  
Gln Glu Glu Lys Leu Lys Glu Met Ala Lys Arg Glu Gly Leu Trp Asn  
65 70 75 80  
Leu Phe Val Pro Ser Phe Asp Gln Leu Phe Gly Glu Gly Leu Thr Asn  
85 90 95



Leu Glu Tyr Gly Tyr Leu Cys Glu Ile Met Gly Arg Ser Val Trp Ala  
 100 105 110  
 Pro Gln Val Phe Asn Cys Gly Ala Pro Asp Thr Gly Asn Met Glu Val  
 115 120 125  
 Ile Leu Arg Tyr Gly Asn Lys Glu Gln Ile Ser Glu Trp Leu Ile Pro  
 130 135 140  
 Leu Leu Glu Gly Arg Ile Arg Ser Gly Phe Ala Met Thr Glu Pro Gln  
 145 150 155 160  
 Val Ala Ser Ser Asp Ala Thr Asn Ile Glu Cys Ser Ile Arg Arg Gln  
 165 170 175  
 Gly Asp Ser Tyr Val Ile Asn Gly Thr Lys Trp Trp Thr Ser Gly Ala  
 180 185 190  
 Met Asp Pro Arg Cys Arg Val Leu Ile Leu Met Gly Lys Thr Asp Phe  
 195 200 205  
 Asn Ala Pro Lys His Lys Gln Gln Ser Met Ile Leu Val Asp Met Arg  
 210 215 220  
 Thr Pro Gly Ile Ser Val Lys Arg Pro Leu Thr Val Phe Gly Phe Asp  
 225 230 235 240  
 Asp Ala Pro His Gly His Ala Glu Ile Ser Phe Glu Asn Val Val Val  
 245 250 255  
 Pro Ala Lys Asn Ile Leu Leu Gly Glu Gly Arg Gly Phe Glu Ile Ala  
 260 265 270  
 Gln-Gly Arg Leu Gly Pro Gly Arg Leu His His Cys Met Arg Leu Ile  
 275 280 285  
 Gly Ala Ala Glu Arg Gly Met Glu Leu Met Ala Gln Arg Ala Leu Ser  
 290 295 300  
 Arg Lys Thr Phe Gly Lys Phe Ile Ala Gln His Gly Ser Phe Val Ser  
 305 310 315 320  
 Asp Leu Ala Lys Leu Arg Val Glu Leu Glu Gly Thr Arg Leu Leu Val  
 325 330 335  
 Leu Glu Ala Ala Asp His Leu Asp Lys Phe Gly Asn Lys Lys Ala Arg  
 340 345 350  
 Gly Ile Leu Ala Met Ala Lys Val Ala Ala Pro Asn Met Ala Leu Lys  
 355 360 365  
 Val Leu Asp Thr Ala Ile Gln Val His Gly Ala Ala Gly Val Ser Ser  
 370 375 380  
 Asp Thr Val Leu Ala His Leu Trp Ala Thr Ala Arg Thr Leu Arg Ile  
 385 390 395 400

Ala Asp Gly Pro Asp Glu Val His Leu Gly Thr Ile Gly Lys Leu Glu  
405 410 415

Leu Gln Arg Ala Ser Lys Leu  
420

<210> 101  
<211> 147  
<212> PRT  
<213> Homo sapiens

<400> 101  
Gly Lys Gly Phe Lys Tyr Ala Met Lys Glu Leu Asp Met Glu Arg Leu  
1 5 10 15

Val Ile Ala Ala Gln Ala Leu Gly Ile Ala Gln Gly Ala Leu Asp Glu  
20 25 30

Ala Ile Pro Tyr Ala Lys Gln Arg Lys Gln Phe Gly Lys Pro Leu Ala  
35 40 45

His Phe Gln Leu Ile Gln Phe Lys Leu Ala Asp Met Ala Thr Lys Leu  
50 55 60

Glu Ala Ala Arg Leu Leu Leu Tyr Arg Ala Ala Trp Leu Ala Asp Arg  
65 70 75 80

Gly Arg Pro Thr Ser Lys Glu Ala Ala Met Ala Lys Leu Phe Ala Ser  
85 90 95

Glu Ala Ala Met Gln Val Ala Asp Asp Ala Val Gln Ile Leu Gly Gly  
100 105 110

Val Gly Tyr Thr Asn Asp Tyr Pro Val Glu Arg Phe Tyr Arg Asp Ala  
115 120 125

Lys Ile Thr Gln Ile Tyr Glu Gly Thr Ser Glu Ile Gln Arg Leu Val  
130 135 140

Ile Ala Arg  
145

<210> 102  
<211> 101  
<212> PRT  
<213> Homo sapiens

<400> 102  
Ala Leu Thr Glu Pro Gly Ala Gly Ser Asp Val Gly Ser Ile Lys Thr  
1 5 10 15

Thr Ala Glu Arg Lys Gly Asp Asp Tyr Ile Leu Asn Gly Ser Lys Met  
20 25 30

Trp Ile Thr Asn Gly Gly Gln Ala Asp Trp Tyr Ile Val Leu Ala Val  
 35 40 45  
 Thr Asp Pro Ala Pro Gly Lys Lys Gly Ile Thr Ala Phe Leu Val Glu  
 50 55 60  
 Lys Asp Thr Pro Gly Phe His Ile Gly Lys Lys Glu Asp Lys Leu Gly  
 65 70 75 80  
 Leu Arg Ser Ser Asp Thr Cys Glu Leu Ile Phe Glu Asp Val Arg Val  
 85 90 95  
 Pro Glu Ser Asn Ile  
 100

<210> 103  
 <211> 22  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: PCR Primer  
 Sequence

<400> 103  
 gaggtctctt ccagtaacat ca

22

<210> 104  
 <211> 26  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: PCR Primer  
 Sequence

<400> 104  
 actctccttg tcctctgagg cgctct

26

<210> 105  
 <211> 22  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: PCR Primer  
 Sequence

<400> 105  
 gcagtttggt tggttggttt ac

22

<210> 106  
 <211> 22

<212> DNA  
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer  
Sequence

<400> 106  
catagccctg tctcaagtct tg

22

<210> 107  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer  
Sequence

<400> 107  
ttccatctct tcagcaaata ctctca

26

<210> 108  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer  
Sequence

<400> 108  
actcttccga catcacaaga aa

22

<210> 109  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PCR Primer  
Sequence

<400> 109  
tgagaatcag atccatgaag ct

22

<210> 110  
<211> 26  
<212> DNA  
<213> Artificial Sequence

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<223> Description of Artificial Sequence: PCR Primer

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